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Aak80342 Human imm
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Aak80349 Human imm
Aak805784 Human imm
Aak80505 Human imm
Aak86605 Human imm
Aak66230 Human imm
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Aak66231 Human gen
Aai67784 Nucleotid
Acn4346 Human DNA
Aas31538 Human pol
Abd66856 Human DNA
Abd44045 Genomic DNA
Abd44045 Genomic DNA
Abd464045 Human DNA
Add261143 Human DNA
Add267038 Human ner
Add26830 Human ner
Add26830 Human syn
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Abx61804 Genomic DNA
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Add2767 Human syn
Abd33219 Human can
Abz35015 Human gen
Adz13610 Human can
Adq17729 Human sof
Abd32888 Human can
Add13739 Osteoarth
Adq19948 Human sof
Adq59197 MSI-H car
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ADQ62941
ID ADQ66
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cytostatic; gene therapy; diagnostic marker; morbid state; osteoporosis;
neurological disease; Alzheimer's disease; Parkinson's disease; dementia;
    21-JAN-2003; 2003JP-00102206
09-MAY-2003; 2003JP-00131392
                                               21-JAN-2004;
                                                                                                                                                                                                                                                                                                               ADQ62941;
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                                                                              28-JUL-2004.
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ADL13748
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ADU404593
ABC83558
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Add36495 Human aut
Ab164403 Stomach c
Ab167239 Thyroid c
Add11748 Osteoarth
Aca61395 Novel human kin
Add640593 Human kin
Add640593 Human kin
Add640593 Human gen
Acn44754 Human gen
Acn44754 Human gen
Abd32843 Human can
Aak65129 Human imm
Aak65129 Human imm
Aak65129 Human imm
Aak65129 Human imm
Abv49425 Human ova
Aak83454 Human pro
Aak83454 Human ova
Add67899 Human imm
Aaa16012 Human ova
Add174275 Human ova
Add174276 Human ova
Add174272 Human ova
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Add139513 Human ova
Add139513 Human ova
Acn91099 Breast ca
Ac139512 Human con
Aal139513 Human con
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Aal139514 Human ova
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Aah10527 Human con
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Best Local Similarity
Matches 3122; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel 2495 cDNA, useful for treating osteoporosis, neurological diseases, Alzheimer's diseases, Parkinson's diseases, dementia and various cancers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Isogai T, 'Yamamoto J,
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                                                                                                                                               GTGGGCGGGTCCTAGGAAACCCTACCCGGCCCCTTGGCAGCGCCTAAGGCGGAGCGCG
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           AGTGCGCCTGGGGAGGATGGACGAGGGAGCGGGGGACCGCTAACGGGGCTCCCTCTGCGC
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                                           <u> AGCTTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGG</u>
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Nagai K, Irie R;
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Pred. No. 0;
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This invention relates to novel, isolated full length human cDNA molecules and the encoded proteins thereof. Specifically, it refers to cDNA clones obtained by an oligo-capping method, where none of these clones are identical to any known human mRNAs. The present invention describes an immunoassay to identify agomNNAs and antagonists, as well antibodies, antisense molecules and siRNAs that can all be used to bind to and modulate expression of the cDNA molecules. As such, these molecules are useful for diagnostic markers or theraparities.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene; ss; human; oligo-capping method; diagnostic marker; geosteoporosis; neurological disease; Alzheimer's disease; Parkinson's disease; dementia; short memory; cancer; sense or motor function; emotional reaction; fear response; osteopathic; neuroprotective; nootropic; antiparkinsonian; osteopathic; neuroprotective; nootropic;
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                                                                                                      ARKS4951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM62170 to AAM91921. (I) have cytostatic CC activity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I), by inserting the CC protein. (I) proteins and polynucleotides may be used to prevent, (CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic decils. AAK64703 to AAK687694 represent human immune/haematopoietic antigen genomic complement sequences used in the exemplification of the present invention. AAK54942 to AAK54950 and AAM62169 represent sequences used in the exemplification of the present invention.
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O O O O O O O O O O O O O O O O O O O	14-SEP-2000; 2000US-023; 14-SEP-2000; 2000US-023; 14-SEP-2000; 2000US-023; 14-SEP-2000; 2000US-023; 14-SEP-2000; 2000US-023; 14-SEP-2000; 2000US-023; 21-SEP-2000; 2000US-023; 25-SEP-2000; 2000US-023; 25-SEP-2000; 2000US-023; 26-SEP-2000; 2000US-023; 27-SEP-2000; 2000US-023; 29-SEP-2000; 2000US-023; 02-OCT-2000; 2000US-023; 02-OCT-2000; 2000US-023; 02-OCT-2000; 2000US-023; 02-OCT-2000; 2000US-023;

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Best Local
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06-DEC-2000; 2000US-0251879P.
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08-DEC-2000; 2000US-0251869P.
08-DEC-2000; 2000US-0251990P.
11-DEC-2000; 2000US-0251990P.
11-DEC-2000; 2000US-0254997P.
05-JAN-2001; 2001US-0259678P.
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AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polymucleotides may be used in the prevention, diagnosis an treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I)
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GAGGTTTTGAGCCAATCAGCTCTGAGACTGGGTTAGAATGTAACAGCTTTAACTTGGGAT
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                                                                                           The invention relates to isolated polynucleotide (I) and polypeptide (II) creation (PCR) primers, oligomers, and for chromosome and gene mapping, come and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed comes. (I) is useful in gene therapy techniques to restore normal comestivity of (II) or to treat disease states involving (II). (II) is complement. (II) and its binding partners are useful in medical imaging complement. (II) and its binding partners are useful in medical imaging complement. (II) and (II) are useful for treating an indication in complement. (II) and (II) are useful for treating disorders complement. (II) and (II) are useful for treating disorders completide and polynucleotide sequences have applications in completide and polynucleotide sequences have applications in composition, forensics, gene mapping, identification of mutations cresponsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and composition of appear in the printed specification, but was obtained in complete contaction format disposition of the invention. Note: The sequence data for this cated the contact of the invention of the sequence data for this contaction format disposition of the printed specification, but was obtained in contaction of the printed specification of the sequence.
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23-AUG-2000;
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                                                                     ftp.wipo.int/pub/published_pct_sequences
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)B; ABG29541.
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RESULT 7
ACH97504/c
ID ACH87504 standard; DNA; 708 BI
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AC ACH87504;
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DT 29-JUL-2004 (first entry)
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                                                                                                                    GCTGAGCTGA 850
                                                                                                                                           GCTGAGCTGA 1565
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99.8%;
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Pred. No. 0;
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Human; probe; ss; gene expression; single exon probe; microarray; alternative splicing event; genomic alteration.
                                                                                                                                                                                                   Human
US2003194704-A1.
                                                     Homo sapiens
                                                                                                                                                                                              genome derived single exon probe #20699
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03-APR-2002; 2002US-00029386 03-APR-2002; 2002US-00029386 (PENN/) PENN S G. (RANK/) RANK D R. (HANZ/) HANZEL D K. Rank DR,

New human genome-derived single exon nucleic acid probes useful for human gene expression analysis, for identifying or characterizing alternative splicing events, for assessing genomic alterations or as tools for surveying tissues.

Claim 1; SEQ ID NO 20699; 80pp; English

CC expression, comprising any of the 27,400 fully defined nucleotide consequences in the specification, or their complements or fragments, and concluding at least 8 amino acids of any of the 6888 amino acid sequences (fully defined in the specification. The probe is a single exon probe that compressed in human cells or rissues. Also included are a spatially-cardinessed in human cells or rissues. Also included are a spatially-cardinessed in human cells or rissues. Also included are a spatially-cardinessed in comprising a plurality of single exon nucleic acid gene expression (comprising a plurality of single exon nucleic acid molecules card probes is a single exon nucleic acid molecules card probes is separately cardinessed in comprising a plurality of probes is separately cardinessed in comprising a plurality of probes is separately cardinessed in comprising a plurality of probes is separately cardinessed in comprising at least gene expression, an excit comprising the single exon nucleic acid molecules contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid contiguous amino acids of any of the above- mentioned amino acid substitutions), an acid substitutions of sequences (optionally with conservative amino acid substitutions), an acid substitutions of sequences of selling and/or licensing single exon probes or microarrays to constitute a proper subscription, and a computer-readable conscituted above. The probe methods and apparatus are useful in gene expression analysis. The probes and apparatus are useful in gene capression analysis. The probes and apparatus are useful in gene capression analysis. The probes and apparatus are useful in gene capression analysis. The probes and apparatus are useful in gene capression of a single exon microarrays. Consider the probes are used in identifying and characterising gross considered and constructions, in detecting and characterising g

Sequence 708 BP; 104 A; 279 C; 231 G; 94 T; 0 U; 0

Query Match Best Local Sim: Matches 707; Similarity Conservative 21.0%; 99.9%; <u>,</u> Score 657; Pred. No. 2. Mismatches DB 12; 2.9e-296; Length 708 Indels <u>.</u> Gaps

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RESULT 8
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ID ACH73793 standard;
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                                                                                                             US2003194704-A1
                                                                                                                                                           Human; probe; ss; gene expression; single exon probe; microarray; alternative splicing event; genomic alteration.
                                                                                                                                                                                               Human genome derived single exon probe #6988.
                                                                                                                                                                                                                           29-JUL-2004
(PENN/)
                                    03-APR-2002; 2002US-00029386
                                                            03-APR-2002; 2002US-00029386
                                                                                    16-OCT-2003
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New human genome-derived single exon nucleic acid probes useful for human gene expression analysis, for identifying or characterizing alternative splicing events, for assessing genomic alterations or as tools for
                                                                                                                                                                                       surveying tissues.
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                                                                                                                                                                                                                       Rank DR,
                                                                                                                                                                                                                        Hanzel
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Claim 15; SEQ ID NO 6988; 80pp; English.

CC measuring human gene expression, a vector comprising the single exon CC probe cited above, an ORF-encoded peptide comprising the single exon CC contiguous amino acids of any of the above- mentioned amino acid sequences (optionally with conservative amino acid substitutions), an CC isolated antibody that binds specifically to a peptide cited above, CC methods of selling and/or licensing single exon probes or microarrays to a customer desiring to measure gene expression, a method of providing CC human gene expression data by subscription, and a computer-readable ceted above. The probe, methods and apparatus are useful in gene capression analysis. The probes may be used as tools for surveying tissues to detect the presence of expressed messages that contain their ceted above. The probes are used in identifying and characterising calternative splicing events, in detecting and characterising calterations in the genomic locus that includes their exon, in assessing smaller genomic alterations, in priming the synthesis of nucleic acids, or in expressing the ORF-encoded peptide. The present sequence is a human constructed in the invention. Note: The sequence data for this captain the corm part of the printed specification, but was obtained consequence. Attail of the printed specification, but was obtained consequence. The miles of the printed specification, but was obtained consequence. The invention relates to a nucleic acid probe for measuring human gene expression, comprising any of the 27,400 fully defined nucleotide sequences in the specification, or their complements or fragments, and encoding at least 8 amino acids of any of the 6888 amino acid sequences fully defined in the specification. The probe is a single exon probe that hybridises under high stringency conditions to a nucleic acid molecule expressed in human cells or tissues. Also included are a spatially-addressable set of single exon nucleic acid probes for measuring human gene expression (comprising a plurality of single exon nucleic acid probes cited above, where each of the plurality of probes is separately and addressably isolatable or amplifiable from the plurality), a single exon microarray for measuring human gene expression, a method of

Sequence 524 BP; 61 A; 212 C; 162 G; 89 T; 0 U; 0 Other;

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Best Local (
                     724
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                                                                                                                                                                                                                                                                                                  484 CTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTCAGC
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CCGAGCGAGTCACGGACCATGAAGAGCGTTCGTGCCGCGGGCCCAAGGCCGGGATGGGG
                                                                              CCGTCCGCAGAGGCGCACGTCGAGGGTCCCCGGGCGCTCCCGTGGACGTTGGCGGTAGCG
                                                                                                                         GCGCCTGGGGAGGATGGACGAGGGAGCGGGGGACCGCTAACGGGGCTCCCTCTGCGCGCCC
                                                                                                                                                                                                                                 TTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGGAGT
                                                        CCGTCCGCAGAGGCGCACGTCGAGGGTCCCGGGGCGGCTCCGTGGACGTTGGCGGTAGCG
                                                                                                                                                            GCGCCTGGGGAGGATGGACGAGGGAGCGGGGACCGCTAACGGGGCTCCCTCTGCGCGCCC
                                                                                                                                                                                              TTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGGAGT
                                                                                                                                                                                                                                                                  CTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTCAGC
                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                     16.8%; Score 524; DB 12;
100.0%; Pred. No. 4.3e-234;
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                       783
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NACK2785;  AAK62785;  AAK62785;  AAK62785;  O6-NOV-2001 (first entry)  Human immune/haematopoietic antigen encoding cDNA SEQ ID NO:7845.  Human; immune/haematopoietic; immune/haematopoietic antigen; cancer; cyrostatic; gene therapy; vaccine; metastasis; ss.  Homo sepiens.  W0200157182-A2.  O9-AUG-2001. 2001W-US001354.  17-JAN-2001; 2001W-US001354.  31-JAN-2000; 2000US-01906SP, 24-FEB-2000; 2000US-01906SP, 24-FEB-2000; 2000US-01907AP, 24-FEB-2000P, 24-FEB-2000P, 24-FEB-2000P, 24-FEB-2000P, 24-FEB-2000P, 2	284 CCGAGCGAGTCACGGACCATGAAGAGCGTTCGTGCCGCGGGCCCAAGGCCGGGATGGGG 225  784 GTTAGCCACATCCTGCCGCGCTGAGGGGAGGCTAACGGGCCGGGCCGGGCCGAGC 843
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2000US-0231968P. 2000US-023239P. 2000US-023239P. 2000US-0233401P. 2000US-0233401P. 2000US-0233401P. 2000US-02334054P. 2000US-0234274P. 2000US-0234998P. 2000US-0234936P. 2000US-0235834P. 2000US-0235836P. 2000US-02353636P. 2000US-0235369P. 2000US-02353779. 2000US-02353799. 2000US-02353799. 2000US-02353799. 2000US-0237039P. 2000US-0237039P. 2000US-0237039P. 2000US-0245675P. 2000US-024478P. 2000US-0244675P. 2000US-0244675P. 2000US-0244679P. 2000US-0246774P. 2000US-0246774P. 2000US-0246774P. 2000US-0246774P. 2000US-0246774P. 2000US-0246774P. 2000US-0246774P. 2000US-0246779P. 2000US-0246778P. 2000US-0246778P. 2000US-0246779P. 2000US-0246673P. 2000US-0246523P. 2000US-0246523P. 2000US-0246533P. 2000US-0246533P. 2000US-0246533P. 2000US-0246533P. 2000US-0246533P. 2000US-0246533P. 2000US-0246611P.	002333333333333333333333333333333333333

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                                                                                                                                                                                    CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cattivity, and can be used in gene therapy and vaccine production. (I) CC cattivity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I), by inserting the CC nucleic acids into a host cell and culturing the cell to express the CC protein. (I) proteins and polynucleotides may be used to prevent, CC diagnose and treat immune/haematopoietic acids; especially CC cancers and cancer metastases of haematopoietic acityen genomic CC sequences from the present invention. AAK54942 to AAK54950 and AAM82169 cc represent sequences used in the exemplification of the present invention.
                                                                                                   Query Match
Best Local Simi
Matches 624;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
06-DEC-2000;
06-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
                                                                                                                                                              Sequence 973 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis.
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17-NOV-2000;
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17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC
                                                                                                                   Local Similarity
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                  2018
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TTCCCGGACAAGAAAATTGCAATCAAATGTCAGCAGCTTTTATTACCTTAATCTTTCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 7845; 3071pp + Sequence Listing; English.
                                            Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000US-0250391P.
2000US-0251030P.
2000US-0251988P.
2000US-0256719P.
2000US-0251879P.
2000US-0251868P.
2000US-0251868P.
2000US-0251869P.
2000US-0251990P.
2000US-0251990P.
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2000US-0249211P.
2000US-0249211P.
2000US-0249211P.
2000US-0249211P.
2000US-0249214P.
2000US-0249214P.
2000US-0249214P.
2000US-0249214P.
2000US-024924P.
2000US-024924P.
2000US-024924P.
2000US-024926P.
2000US-024926P.
2000US-024929P.
2000US-0249300P.
2000US-0249300P.
                                                                                                       Conservative
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                                                                                                                                                              255 A; 253 C; 226 G; 234 T; 0 U; 5 Other;
                                                                                                                  16.1%;
                                                                                                   Score 504; DB 4; Pred. No. 9.3e-225;
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RESULT 10
AAK83430/c
ID AAK834
AC AAK834
AC AAK834
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DT 07-NOV
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Human
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24-FEB-2000
02-MAR-2000
16-MAR-2000
17-MAR-2000
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19-MAY-2000
07-JUN-2000
28-JUN-2000
07-JUN-2000
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07-JUN-2000
07-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; immune; haematopoietic; immune/haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immune/haematopoietic antigen genomic sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic; gene therapy; vaccine; metastasis;
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AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis an treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting th nucleic acids into a host cell and culturing the cell to express the
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01-DEC-2000

06-DEC-2000

08-DEC-2000

AK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis are
                                                                                                                 Nucleic
                                                                                                                                                                        Rosen
                                                                                                                                             2001-483426/52
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                                                                                                 : acids encoding
for preventing,
                                                                                                                                                                                                  HUMAN GENOME
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                                                                     NO 38239; 3071pp +
                                                                                                                                                                                                    SCI
                                                                                                 human immune/hematopoietic diagnosing and/or treating
                                                                                                                                                                      Ruben
                                                                                                                                                                        S
                                                                     Sequence
                                                                     Listing;
                                                                                                 antigen polypeptides, cancers and metastasis.
                                                                     English
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RESULT 12
AAK83426/c
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Best Local S
Matches 471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the protein. (I) proteins and polynucleotides may be used to produce the secreted (I), by inserting the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK59422 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 476 BP; 59 A; 190 C; 157 G; 70 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased
                                                                                                                                                                                                                  07-NOV-2001
                                                                                                                                              Human; immune; haematopoietic; immune/haematopoietic antigen; cancer:
                                                                                                                                                                               Human immune/haematopoietic antigen genomic sequence
                                                                                                                                                                                                                                                                                 AAK83426
                              09-AUG-2001.
                                                               WO200157182-A2
                                                                                                                             cytostatic;
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ilarity 100.0%;
Conservative
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                                                                                                                                 therapy; vaccine; metastasis;
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Pred. No. 2.5e-209;
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prevention, diagnosis and

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16-MAR-2000
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Nucleic acids encoding useful for preventing,
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  cancers
  polypeptides, and metastasis.
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Best Local S
Matches 427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 476 BP; 58 A; 191 C; 157 G; 70 T; 0 U; 0 Other;
           Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 38238; 3071pp + Sequence Listing; English
                                                DNA encoding novel human diagnostic protein #8312.
                                                                         13-FEB-2002
                                                                                                 AAS72508
                                                                                                                       AAS72508 standard; cDNA; 1349
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                427;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        acid sequences
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                                                                                                                                                                                                                                                                                                                                                                                                          CAACAAGACGACTGCGTGCTACCACCACCTGGTGCTGACCGTCGGTGGCTCGGCGGACTC
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                                                                                                                                                                                                                                                         CGAGTTCGAGCGGCTCTGGGTGGCCTTCTCGGGCTGCCTGGACCTGCTGGAAGCGGACAT 1135
                                                                                                                                                                                                                                                                                   AAK64702 encode the human immune/haematopoietic antigen
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                                                                         (first
                                                                                                                                                                                     50
                                                                         entry)
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                                                                                                                        βP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 476;
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The invention relates to isolated polynucleotide (I) and polypeptide (II) creation (PCR) primers, oligomers, and for chromosome and gene mapping, cc reaction (PCR) primers, oligomers, and for chromosome and gene mapping, cc and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal cc useful for generating antibodies against it, detecting or quantitating a collection of till and its binding partners are useful in medical imaging cc supplement. (II) and its binding partners are useful in medical imaging cof sites expressing (II). (I) and (II) are useful for treating disorders cinvolving aberrant protein expression or biological activity. The cc involving aberrant protein expression or biological activity. The cc diagnostics, forensics, gene mapping, identification of mutations composible for genetic disorders or other traits to assess biodiversity and no produce other types of data and products dependent on DNA and continuous according sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention. Note: The sequence data for this categories of continuous directly from WIPO at the continuous continuous contact directly from WIPO at the continuous continuous contact directly from WIPO at the contact of 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated polynucleotide and encoded polypeptides, useful diagnostics, forensics, gene mapping, identification of mutat responsible for genetic disorders or other traits and to asse biodiversity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAR-2000; 2000US-00540217.
23-AUG-2000; 2000US-00649167.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ftp.wipo.int/pub/published_pct_sequences
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C 1142
                                                                                                                                                       CGAGCGGCTCTGGGTGGCCTTCCTCGGGCTGCCTGGACCTGCTGGAAGCGGACATGCGACG 114:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   CCTGCGGCAGGAGCTGCAAAAGACGCGCCAGAAGGCGCAGGAGCTGGCGGTGTCCACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCTGCGGCAGGAGCTGCAAAAAGACGCGCCAGAAGGCGCAGGAGCTGGCGGTGTCCACCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GACGACTGCGTGCTACCACCACCTGGTGCTGACCGTCGGTGGCTCGGCGGACTCGCAGAA
                                                                                           CGAGCGGCTCTGGGTGGCCTTCTCGGGCTGCCTGGACCTGCTGGAAGCGGACATGCGACG
                                                                                                                                                                                                                                                                         CGCCCGGCTGACTGCTGCTGCGCGACCGGGGCCTGGCCGACGACGAGCGCCGAGTT
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RESULT 14 ABN50582

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                                                                                                                                                                                                     The present invention describes oligonucleotide libraries for detecting CC messenger RNAs that populate a (sub-)transcriptome, where the (sub-CC )transcriptome comprises messenger RNAs transcribed from multiple CC transcription units that populate a genome. The library comprises several CC oligonucleotides, each capable of hybridising selectively to a set of CC messenger RNAs transcribed from a given transcription unit of the genome, CC which encodes one or more messenger RNA splice variants. The CC oligonucleotide libraries are useful for detecting mRNAs from a CC discounties and publicatively or CC detecting RNA transcripts and splice variants of numan or animal CC transcriptomes. The libraries may also be used as specialised mini CC libraries to detect transcripts of a sub-transcriptome under a particular biological or pathological state, and so allowing the detection of tissue cand pathology-specific genes such as those genes only expressed in CC developmental specific genes such as those genes only expressed in CC specific tissue under a specific pathological condition, to detect can pathological or pathological or pathological or pathological or pathological or specific pathological condition, to detect condition to detect and pathological or pathologi
                                                                                     Best Loc
Matches
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Best Local Similarity
                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription unit genome, useful for detecting tissue-, pathology-, and developmental-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; mouse; rat; splice transcript; detection; RNA transc splice variant; transcriptome; oligonucleotide library; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; SEQ ID NO 23330; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  specific genes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human spliced transcript detection oligonucleotide SEQ ID NO:23330.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-JUL-2000; 2000US-0221607P
02-MAY-2001; 2001US-0287724P
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                          2002-257383/30
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  BP; 11 A; 14 C; 16 G; 19 T; 0 U; 0 Other;
                                                                                     Conservative
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RESULT 15
ABK83571/c
               CC (GCA), by detecting the level of expression of gene(s) (GS) identified by CC (DNA chip analysis as given in the specification, and comparing the CC expression level to an expression level in an unactivated GC, where CC expression of at less one of Gs is indicative of GCA. Also included are CC (condulating (M2) GA by contacting GC with an agent that alters the CC expression of at less tone gene in Gs; (2) screening (M3) for an agent CC expression of at less tone gene in Gs; (2) screening (M3) for an agent CC expression of at less on a subject, exposure of a subject to a pathogen or sterile inflammatory disease using the gene expression CC pathogen or sterile inflammatory disease, by detecting the level of CC expression in a sample of the tissue of gene(s) from Gs, where the level of CC expression in a subject, exposure of a subject to a pathogen or sterile inflammatory disease, by detecting the level of CC expression in a subject, exposure of inflammation; (4) treating CC (M5) an inflammatory disease, by contacting a tissue having inflammation with an CC is useful for detecting GCA; M2 is useful for modulating GA; M3 is useful for screening an agent capable of gene(s) from Gs in the tissue, an allergic exposure of a subject to a pathogen or sterile inflammation in a tissue; M1 is useful for detecting GA; M2 is useful for modulating GA; M3 is useful for detecting GA; M3 is useful for detecting GA; M3 is useful for detecting an inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammation (especially chronic) in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease (e.g. general infection, viral infection, parasitic infection, protozoal infection, conditions. The present sequence represents a gene differentially conditions. The present sequence are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; 88; granulocytic cell; DWM CHAP, DURCOLO Infection; vital infection; parasitic infection; protozoal infection; fungal infection; sterile inflammatory disease; psoriasis; rheumatoid arthritis; glomerulomephritis; asthma; thrombosis; cardiac reperfusion injury; renal reperfusion injury; ARDS; adult respiratory distress syndrome; inflammatory bowel disease; adult respiratory distress syndrome; inflammatory disease; Crohn's disease; ulcerative colitis; periodontal disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detecting granulocyte activation by detecting differential expression of genes associated with granulocyte activation, which serves as diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 142; 114pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to detecting (M1) granulocyte (GC) activation
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Best Local
                                                                                                                    The invention relates to a method of determining susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain comprising identifying whether the individual has at least one polymorphism in a polynucleotide encoding at least one of the protein listed in the specification. The methods, composition and agent are useful for modulating the susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain that is associated with a disease, preferably osteoarthritis. The cell line and the non-human animal are useful for screening for an agent for diagnosing an individual having susceptibility to joint space narrowing and/or osteophyte development and/or joint pain. This sequence corresponds to the polynucleotide encoding a protein listed in the specification. (Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                               Determining susceptibility of an individual to joint space narrowing, ostsophyte development and/or joint pain comprises identifying whether the individual has at least one polymorphism in a polynucleotide encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ds; gene; osteopathic; antiinflammatory; antiarthritic; gene therapy; joint space narrowing; osteophyte development; joint pain; osteoarthritis; SNP; single nucleotide polymorphism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jones KA,
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                                                                      Sequence 175737 BP; 41985 A; 43790 C; 42407 G; 47555 T; 0 U; 0 Other;
                                                                                                       ftp.wipo.int/pub/published_pct_sequences).
                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 128; 297pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (INCY-) INCYTE GENOMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  53;
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                    Similarity
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 ilarity 100.0%; I Conservative 0;
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                  1.7%; Score 53;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.7%; Score 53;
100.0%; Pred. No.
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 Mismatches
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                                    DB 10;
                  4.9e-14;
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                                   Length 175737;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                #128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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   0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
   Gaps
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SAXE

AAC03795

AAC03795 standard;

CDNA;

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RESULT 18
AAC03795/c
                                                                                                                              Matches
                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                      which comprises obtaining a first soft tissue sample from an individual and a normal soft tissue sample from the same or different individual, determining the expression of a gene in both samples and comparing the expression in the first soft tissue sample indicates the presence of soft tissue sarcoma. The method of the invention has cytostatic applications and may be useful for detecting soft tissue sarcoma, possibly via gene therapy or vaccine production. The nucleic acid sequences may be useful in diagnostic and screening applications. The current sequence is that of a human soft tissue sarcoma-upregulated DNA of the invention. The current sequence is not shown within the specification per se but was submitted in CD format by the inventor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Early detection of soft tissue sarcoma comprises determining expression of a gene in a first soft tissue sample and a normal soft tissue sample and comparing the gene expression, also useful in treating soft tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human soft tissue sarcoma-upregulated DNA - SEQ ID 1753.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADQ18934 standard; DNA; 175737
                                                                                                                                                                                          Sequence 175737 BP; 41985 A; 43790 C; 42407 G; 47555 T; 0 U; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aziz N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-NOV-2002; 2002US-0429739P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-NOV-2003; 2003WO-US038193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel method for detecting soft tissue sarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and comparing the gene expression,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-441208/41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       soft tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2; SEQ ID NO 1753; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PROT-)
                                                                                                                                             ocal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                47829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2888
                                                                                 2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT
                                                                                                                              53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ginsburg WM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 47777
                                                                TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sarcoma;
                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry
                                                                                                                     1.7%; Sur.
/ 100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              zlotnik A;
                                                                                                                                              Score 53; DB 12;
Pred. No. 4.9e-1
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                                                                                                                               Mismatches
                                                                                                                                             4.9e-14;
                                                                                                                               0
                                                                                                                                                            Length 175737;
                                                                                                                               Indels
                                                                                                                              0
                                                                                                2940
                                                                  47777
                                                                                                                                                                                               Other;
                                                                                                                              Gaps
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RESULT 19
AAS78337/C
ID AAS78
XX AAS78
AC AAS78
XX AAS78
CT 13-FE
XX DNA 6
XX Humar
KW Hood
XX Homo
XX Homo
XX Homo
XX WO201
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                                                        Matches 52;
                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion
                                                                               Human; chromosome mapping; gene mapping; gene therapy; food supplement; medical imaging; diagnostic; genetic of
                                                                                                                                                                 13-FEB-2002
                                                                                                                                                                                                                                    AAS78337 standard; cDNA; 1437 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 381 BP; 73 A; 98 C; 84 G; 123 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dumas Milne Edwards J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-FEB-2000; 2000EP-00200610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene
                 WO200175067-A2
                                                 Homo sapiens
                                                                                                                              DNA encoding novel human diagnostic protein #14141.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is one of a large number of 5' ESTs derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 3793; 71pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; 5' EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human secreted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-OCT-2000
                                                                                                                                                                                                                                                                                                                                         3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGAGAAGACTCTGTCTC 3122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                       107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000-500381/45
                                                                                                                                                                                                                                                                                                                     CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 56
                                                                                                                                                                                                                                                                                                                                                                                      Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry
                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expressed sequence tag; secreted protein; cDNA isolation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0122487P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein 5' EST, SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Duclert A,
                                                                                                                                                                                                                                                                                                                                                                                     ; Score 52; DB 
%; Pred. No. 1.8
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Giordano
                                                                                                                                                                                                                                                                                                                                                                                     1.8e-13;
hes 0;
                                                                                                                                                                                                                                                                                                                                                                                                                        DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3793.
                                                                                                                                                                                                                                                                                                                                                                                                                        Length 381;
                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                disorder; ss
                                                                                                forensic;
                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 20
AAL06207/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cc in diagnostics as expressed sequence tags for identifying expressed cc genes. (I) is useful in gene therapy techniques to restore normal cc activity of (II) or to treat disease states involving (II). (II) is constituted in the sequence of the treat disease states involving (II) is consider the sequence of the sequence of sites expressing (II). (II) and its binding partners are useful in medical imaging consultates expressing (II). (II) and (II) are useful in medical imaging consultates expressing (II). (II) are useful for treating disorders convolving aberrant protein expression or biological activity. The colypeptide and polynucleotide sequences have applications in colypeptide and polynucleotide sequences have applications consisting forensics, gene mapping, identification of mutations consisting for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and camino acid sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention. Note: The sequence data for this patent did not appear in the printed specification, but was obtained in celectronic format directly from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                           Human; reproductive system related antigen; reproductive cancer; gene therapy; ds.
                                                                                                                                                                                                                           21-NOV-2001
                                                                                                                                                                                                                                                                                                     AAL06207 standard; DNA; 9620 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1437 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polymucleotides are also used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAR-2000;
23-AUG-2000;
                                                                                                                                                                                     Human reproductive system related antigen DNA SEQ ID NO: 8895.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 14141; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                biodiversity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-639362/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Drmanac RT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAR-2001; 2001WO-US008631
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity nes 52; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                       3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
                                                                                                                                                                                                                                                                                                                                                                                                   243
                                                                                                                                                                                                                                                                                                                                                                                                   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-00540217.
2000US-00649167.
                                                                                                                                                                                                                           (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      282 A; 398 C; 395 G; 362 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.7%; Score 52;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 1.'
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.7e-13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1437;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                 system disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                   192
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02-AUG-2001 WO200155320-A2

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31-JAN-2000

04-FBB-2000

24-FBB-2000

10-MAR-2000

11-MAR-2000

11-MAR-2000

07-JUN-2000

07-JUN-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000

11-AUG-2000

11-SEP-2000

11-SEP-2000

01-SEP-2000

02-SEP-2000

03-SEP-2000

04-SEP-2000

05-SEP-2000

06-SEP-2000

07-SEP-2000

08-SEP-2000

09-SEP-2000

29-SEP-2000

29-SEP-2000

29-SEP-2000

29-SEP-2000
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2000US-0179065P.
2000US-018464P.
2000US-018466P.
2000US-018466P.
2000US-0198123P.
2000US-029467P.
2000US-0216486P.
2000US-0216486P.
2000US-021649P.
2000US-021649P.
2000US-021649P.
2000US-021649P.
2000US-0217496P.
2000US-0225214P.
2000US-0225214P.
2000US-0225266P.
2000US-0225266P.
2000US-022575P.
2000US-023141P.
2000US-023144P.
2000US-023149PP.
2000US-023499P.
2000US-0235836P.
2000US-023636P.
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    29-SRP-2000
02-OCT-2000)
02-OCT-2000)
02-OCT-2000)
02-OCT-2000)
03-OCT-2000)
03-OCT-2000)
03-OCT-2000)
03-OCT-2000)
03-OCT-2000)
03-OCT-2000)
03-NOV-2000)
03-NOV
   Isolated nucleic acid molecule encoding a reproductive system antigen
                                                                 Rogen
                                                                                              (HUMA-)
                                 2001-465570/50
                                                               ß
                                                               Barash
                                                                                                                         2000US-0236370P.
2000US-023703PP.
2000US-023703PP.
2000US-0237039P.
2000US-0237039P.
2000US-0237049P.
2000US-0237049P.
2000US-0241966P.
2000US-0241808P.
2000US-0241808P.
2000US-0241808P.
2000US-0246471P.
2000US-0246471P.
2000US-0246477P.
2000US-0246477P.
2000US-0246477P.
2000US-0246478P.
2000US-0246478P.
2000US-0246523P.
2000US-0249211P.
2000US-0249211P.
2000US-0249211P.
2000US-0249211P.
2000US-024921P.
2000US-0251869P.
2000US-0251869P.
2000US-0251989P.
2000US-0251999P.
2000US-0251999P.
2000US-0251999P.
2000US-0251999P.
2000US-0251999P.
2000US-0251999P.
2000US-025499P.
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                                                                 Ruben
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RESULT AND STATE OF S
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Best Local S
Matches 52
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The present invention relates to methods for distinguishing between individuals homozygous for and therefore afflicted with Van Buchem's disease, individuals heterozygous for and therefore carriers of Van Buchem's disease and individuals who are not afflicted with Van Buchem's Buchem's disease comprise identifying a large genomic deletion in chromosome 17 at 17q21. The method is useful for identifying individuals who are afflicted with or carriers of diseases associated with one or more genomic
                                                                                                                                                                                                                                  Methods for identifying subjects who are afflicted with or carriers of diseases associated with genomic deletion(s), e.g. Van Buchem's disease, by determining the presence of a deletion in the 92 kb region of human chromosome 17 at 17q21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JUL-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; Van Buchem's disease; genomic deletion; craniotubular hautosomal recessive disorder; chromosome 17; chromosome 17q21; bone dysplasia; 92Kb gene fragment; ds.
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                                                                                                                                                                                      Claim 14; Page 45-72; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JUL-2000; 2000US-0221855P 06-JUL-2001; 2001US-0303386P
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(STRA/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAEHLING HAMPTON K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CAAGATTGTGCCACTGCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/note= "This region is deleted in individuals afflicted
or carriers of Van Buchem's disease"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2586 A; 2358 C;
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Best Local S
Matches 52
                                                                                                 Determining a risk for or presence of altered bone mineral density (e. osteoporosis) in a subject comprises determining the presence or abser of a sclerostin gene region nucleotide polymorphism in a biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        deletion, particularly Van Buchem's disease, which is a rare recessive disorder that results in a bone dysplasia referred craniotubular hypertosis. The present sequence is a 92Kb gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  variation
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RESULT 23
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                presence or absence of at least one sclerostin gene region nucleotide polymorphism in a biological sample from a subject where the presence of at least one polymorphism at a position that corresponds to a non-coding region of the 130320 bp sclerostin gene region (SOST) indicates an increased risk of altered BMD. The composition and methods are useful in determining in a subject a risk for having, or presence of, altered bone fracture or other conditions characterized by decreased or increased bone fracture or other conditions characterized by decreased or increased bone density. These may also be used in identifying agents that may be used for treating the above diseases, disorders or conditions associated with altered BMD. In addition, these may be used for pharmacogenomic purposes, e.g. to stratify patient populations according to suitability of a particular therapeutic agent for use in the population. This sequence corresponds to the human sclerostin gene region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SMRT inhibitor; cytostatic; antiinflammatory; antiarthritic; antirheumatic; antisense therapy; inflammatory disorder; rheumatoid arthritis; hyperproliferative disorder; cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMRT; silencing mediator for retinoid and thyroid hormone action;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human SMRT partial genomic DNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of altered bone mineral density (BMD) in a subject by determining
                                                                                                                                                                     (silencing mediator for retinoi treating animal having disease
                                                                                                                                                                                        Novel antisense compound targeted to nucleic acid encoding SMRT (silencing mediator for retinoid and thyroid hormone action), u
                                                                                                                                                                                                                                           GENBANK; NT_009459.
                                                                                                                                                                                                                                                                                               Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                   17-JUN-2002; 2002US-00174014.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              breast cancer; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG86300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG86300 standard; DNA; 220756
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                                                                                                                                                                                                                                                               2004-082184/08
                                                                                                                                                                                                                                                                                                                                  ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                               Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO:14.
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                                                                                                                                                                                         useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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targeted to a nucleic acid molecule encoding SMRT (silencing mediator for retinoid and thyroid hormone action), where (I) specifically hybridises with the nucleic acid molecule encoding SMRT and inhibits expression of SMRT. (I) specifically hybridises with at least 8-nucleobase portion of preferred target region on nucleic acid molecule encoding SMRT. Also

The present invention describes a compound (I) 8-50 nucleobases in length

for

Example 15;

SEQ ID NO 14; 260pp;

English.

associated with SMRT such as cancer,

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The present invention relates to novel DNA and protein sequences which are associated with carcinomas. The sequences are useful for: (1) for screening drug candidates; (11) for screening of bloactive agent capable of bioactive agent capable of bioactive agent capable of modulating the activity of CAP; (11) for screening of bloactive agent capable of modulating the activity of CAP; (1v) for carcinoma; (vi) for inhibiting the activity of CAP; (vi) for treating carcinoma; (vi) for neutralizing the effect of CAP; (vii) for treating Carcinoma; (viii) for neutralizing the effect of CAP; (ix) as a biochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of CA genes are useful as DNA vaccines and the CAP are useful as markers of CAP carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 52;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cytostatic; carcinoma; lymphoma; cancer; human; gene;
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                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 652; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                              Recombinant nucleic acid useful for diagnosis and treatment of carcinoma comprises a nucleotide sequence.
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Pred. No.
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                                                                                        isolation and characterisation of the DNA and protein sequences of the invention. The breast and ovarian cancer associated DNA, protein, agonist or antagonist sequences exhibit cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antibacterial; antiinflammatory; antiulcer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic and cardiant activity. The polynucleotide and protein sequences are used in the diagnosis of cancer, particularly breast and ovarian cancer. The nucleic acid sequences, proteins, agonists and agonists may also be used in the diagnosis, prevention and treatment of immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, militing scolaris repureroid arthritis and ulcerative colitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; breast cancer; ovarian cancer; cytostatic; immunosuppressive; nootropic; neurpprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antivicer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic; cardiant; immune disorder; Addison's disease; allergy; autoimmune haemolytic anaemia; autoimmune thyroiditis; diabetes mellitus; Crohn's disease; multiple sclerosis; rheumatoid arthritis; ulcerative colitis; cardiovascular disorder; wound healing; neurological disease; ds.
   disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis, cardiovascular disorders such as myocardial ischaemias; wound healing; neurological diseases such as cerebral anoxia and epilepsy; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequences AAF21614 - AAF22031 represent DNA sequences encoding human proteins AAB58711 - AAB59128. The DNA and protein sequences are associated with breast and ovarian cancer. Included in the invention sequences AAF22032 - AAF22040 and AAB59129 which are used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New human breast and ovarian cancer associated gene sequences and the polypeptides encoded by these genes, useful in the prevention, treatmund diagnosis of cancer, immune disorders, cardiovascular disorders as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rosen CA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
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Pred. No.
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1.4e-13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention are
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RESULT 26
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                         CC sequences. (I) is useful as hybridisation probes, polymerase chain Creaction (PCR) primers, oligomers, and for chromosome and gene mapping, CC and in recombinant production of (II). The polymucleotides are also used CC in diagnostics as expressed sequence tags for identifying expressed CC genes. (I) is useful in gene therapy techniques to restore normal CC genes. (I) or to treat disease states involving (II). (II) is useful in gene therapy techniques to restore normal CC generating antibodies against it, detecting or quantitating a CC polypeptide in tissue, as molecular weight markers and as a food CC supplement. (II) and its binding partners are useful in medical imaging CC is sites expressing (II). (I) and (II) are useful for treating disorders CC involving aberrant protein expression or biological activity. The CC polypeptide and polymucleotide sequences have applications in CC diagnostics, forensics, gene mapping, identification of mutations companies, forensics, gene mapping, identification of mutations and to produce other types of data and products dependent on DNA and CC amino acid sequences. AAS64197-AAS94564 represent movel human diagnostic coding sequences of the invention. Note: The sequence data for this content did not appear in the printed specification, but was obtained in CC electronic format directly from MIDO at the print of the inventors.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 29534; 103pp; English.
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23-AUG-2000;
             ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to isolated polynucleotide (I) and polypeptide (II)
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                                                                                                                                                                                                                                                                                    detected. (M1) involves identifying nucleic acid sequences comprising comethylated CpG site in promoter-first exon region and that are down-regulated in diseased cells, comparing expression level of nucleic acid sequences with that of demethylated nucleic acid sequences and comparing expression level of nucleic acid sequences exhibiting increase in expression comparing comparing expression of the disease in a subject, which involves determining the degree of comparing of a disease in a subject, which involves determining the presence of the disease in the subject and determining the presence of comparing the obtained from the subject, and determining the presence compared position to, or stage of the disease in the subject based on the degree of methylation; (2) monitoring the onset, progression, or compared for inhibiting a disease in a subject; and (4) a kit (1) useful for diagnosis, prognosis, staging, monitoring, and therapeutic compared for a disease. (M1) is useful for identifying one or more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying nucleic acid sequences as biomarker for disease, by identifying nucleic acid sequences comprising methylated CpG site and down-regulated in diseased cells and comparing its expression level with
                                                                                                                          nucleic acid sequences useful as a biomarker for a disease to be detected, where the nucleic acid sequences are useful for detecting, the presence or stage of a disease such as cancer e.g. colorectal cancer in a subject. The present sequence represents a specifically claimed human genomic sequence for use in the method of the invention. Note - The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Beard C, Burgess C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a method (M1) for identifying one or more nucleic acid sequences useful as a biomarker for a disease to be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    demethylated nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA methylation; biomarker; cancer; gene; ds; SLC26A2.
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                                                                                               sequence data
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ANK; AI025519.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 22; 27pp; English.
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                                                                for this patent is not represented in the printed but was obtained in electronic format from the USI
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Pred. No.
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RESULT 28
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11-JUL-2000;
11-JUL-2000;
11-JUL-2000;
14-JUL-2000;
26-JUL-2000;
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14-AUG-2000;
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14-AUG-2000;
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28-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytostatic; gene therapy; vaccine; metastasis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22051.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2890 AGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
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2000US-0209467P.
2000US-0215135P.
2000US-0216647P.
2000US-0216880P.
2000US-0217486P.
2000US-02174818P.
2000US-022964P.
2000US-022513P.
2000US-0225214P.
2000US-0225214P.
2000US-0225266P.
2000US-0225268P.
2000US-0225268P.
2000US-022575P.
2000US-02259343P.
2000US-02259345P.
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2000US-0198123P.
2000US-0205515P.
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Pred. No.
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RESULT 29
ADX80722/c
ID ADX80722;
XX ADX80722;
AC ADX80722;
XX O5-MAY-20
XX DB Human man
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic contivity, and can be used in gene therapy and vaccine production. (I) CC activity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I), by inserting the CC mucleic acids into a host cell and culturing the cell to express the CC protein. (I) proteins and polynucleotides may be used to prevent, CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic acids. AAK64703 CC sequences from the present inwunne/haematopoietic antigen genomic CC sequences from the present invention. AAK54942 to AAK54950 and AAM82169 CC represent sequences used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 1.6%;
Best Local Similarity 100.0%;
Matches 51; Conservative (
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17-NOV-2000;
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                          Human mannose receptor C type 2 (ENDO180)
                                                                                        05-MAY-2005
                                                                                                                                             ADX80722;
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                                genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 30393;
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                                   DNA.
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01-NOV-2000

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2000US-024677P.
2000US-024677P.
2000US-0246611P.
2000US-024652P.
2000US-0246611P.
2000US-024661P.

melanoma; DNA melanoma; DNA single nucleo single nucleo Homo sapiens.  Key variation variation variation variation variation	variation	variation	variation	variation	variation	variation	variation	variation	variation	variation	VOL TOCHOLL	variation	variation	variation variation	variation variation variation	variation variation variation variation	variation variation variation variation variation variation	variation variation variation variation variation variation	variation variation variation variation variation variation variation variation
nucleotide polymorphism; SNP detection; Cytostatic; gene therapy; SNP; nucleotide polymorphism; gene; ds; chromosome 17.  Location/Qualifiers  224	g= f ndard_name= "Single nucleotide	nucleotide	name= "Single nucleotide	= "Single nucleotide	name= "Single nucleotide	name= "gingle mucleotide	reme=ormAre mucreorine	22	/"crays " /standard_name= "Single nucleotide polymorphism" 19195	<pre>/*tegg= n /*tandard_name= "Single nucleotide polymorphism" 19925</pre>	<pre>/*tag= o /standard_name= "Single nucleotide polymorphism" 10040</pre>	/*tag= p /standard_name= "Single nucleotide polymorphism"	20300	<pre>/*tag= q /standard_name= "Single nucleotide polymorphism"</pre>	q rd_name= "Single nucleotide r rd name= "Single nucleotide	_name= "Single nucleotide polymo _name= "Single nucleotide polymo	d name= "Single nucleotide r r d_name= "Single nucleotide s d_name= "Single nucleotide	_name= "Single nucleotide polymc _name= "Single nucleotide polymc _name= "Single nucleotide polymc _name= "Single nucleotide polymc	_name= "Single nucleotide polymc
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variation variation variation variation variation variation	variation	variation	variation	<b>Z.</b>	Ħ.	3.		variation	variation	variation	variation	variation	variation	variation		variation	variation		variation
w "Single nucleotide x "Single nucleotide x "Single nucleotide y "Single nucleotide ard_name= "Single nucleotide ac ac "Single nucleotide ad	22	/*tag= ae /standard_name= "Single nucleotide polymorphism" 39707	/*teg= af /standard_name= "Single nucleotide polymorphism" 40072	<pre>/*tag= ag /standard_name= "Single nucleotide polymorphism" 41164</pre>	<pre>/*teg= ah /standard_name= "Single nucleotide polymorphism" 41767</pre>	<pre>/*tag= ai /standard_name= "Single nucleotide polymorphism" 42724</pre>	<pre>/*teg= aj /*tag= aj /standard_name= "Single nucleotide polymorphism"</pre>	43139 /*tag= ak /standard name= "Single nucleotide polymorphism"	1	<pre>/standard_name= "Single nucleotide polymorphism" 49720 /*tag= am</pre>	12	#	/standard_name= "Single nucleotide polymorphism" 51853	/"tag= ap /standard_name= "Single nucleotide polymorphism" 51946	<pre>/*tag= aq /standard name= "Single nucleotide polymorphism"</pre>		<pre>/standard_name= "Single nucleotide polymorphism" 59414 /*tag= as</pre>		22

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ACN44754/c
ID ACN447
XX
AC ACN447
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DT 18-NOV
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DT 18-VOV
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Human
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OS Homo s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel method for identifying a subject at risk of melanoma. The method comprises detecting the presence or absence of a copolymorphic variation associated with melanoma, where the presence of the cone or more polymorphic variations is indicative of the subject being at risk of melanoma. The invention further comprises: a method for identifying a polymorphic variation associated with melanoma proximal to an incident polymorphic variation associated with melanoma; an isolated nucleic acid which comprises a portion of or all of a nucleotide sequence comprising fully defined 68400-213300 base pairs sequences (SEO ID NO. 3, 4, 5, 6, and/or 7) given in the specification, and comprises one or more polymorphic variations; an oligonucleotide comprising a nucleotide sequence complementary to a portion of the nucleotide sequence above, where the 3' end of the oligonucleotide to adjacent to a polymorphic variation; a microarray comprising the isolated nucleic acid linked to solid support; an isolated polypeptide encoded by the isolated nucleic acid sequence; genotyping a nucleic acid; a method for identifying a
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                       Matches 51;
                                    WO2003073826-A2
                                                                                                      Cytostatic; carcinoma; lymphoma; cancer; human;
                                                                                                                                        Human genomic sequence hCG37990.
                                                                                                                                                                             18-NOV-2004
                                                                                                                                                                                                                                                ACN44754 standard; DNA; 215221 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a subject at risk of melanoma by detecting presence or absence of a polymorphic variation associated with melanoma, where the presence of polymorphic variations is indicative of the subject being risk of melanoma.
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06-NOV-2003; 2003US-00703789

06-NOV-2003; 2003US-00703817

06-NOV-2003; 2003US-00704513
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   12-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                                                                                                      2890 AGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
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llarity 100.0%;
Conservative (
                                                                                                                                                                             (first entry)
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Pred. No. 4.3e-13;
                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                      gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 68200;
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RESULT 31
AAC24464/c
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Best Local Similarity
Matches 51; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of bioactive agent capable of modulating the activity of CAP; (iv) for a bioactive agent capable of modulating the activity of CAP; (iv) for carcinoma; (vi) for inhibiting the activity of CAP; (vi) for treating carcinoma; (vii) for number of a carcinoma drug; (vii) for reating the effect of CAP; (vi) as a biochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent inspands of the capable patent to basic patent in the capable patent.
New nucleic acid that is a 5' expressed sequence obtaining cDNAs and genomic DNAs that correspond
                                                WPI; 2000-500381/45
                                                                           Dumas Milne Edwards J,
                                                                                                                                        26-FEB-1999;
                                                                                                                                                                      21-FEB-2000; 2000EP-00200610
                                                                                                                                                                                                    06-SEP-2000
                                                                                                                                                                                                                                   EP1033401-A2
                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                           gene therapy;
                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                         Human secreted protein 5' EST, SEQ ID NO: 28539.
                                                                                                                                                                                                                                                                                                                                                                         06-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC24464 standard; cDNA; 255
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 1360; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Recombinant nucleic acid useful for diagnosis and treatment of carcinoma comprises a nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAR-2002; 2002US-00087192
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT 3121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention relates to novel DNA and protein sequences which
                                                                                                                                                                                                                                                                                                           5' EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.6%; So ilarity 100.0%; F Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                         ; expressed sequence
chromosome mapping;
                                                                                                                                         99US-0122487P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for which no sequence data was
                                                                              Duclert
                                                                                                                                                                                                                                                                                              mapping; ss.
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                                                                              Giordano
                                                                                                                                                                                                                                                                                                           secreted protein;
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hes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    published
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 215221;
 tag (5' EST)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cardiovascular; respiratory; gene therapy; secreted protein; chromosome identification; hybrid mapping; gene expression control; immune system disorder; immunodeficiency; Chediak-Higashi syndrome; autoimmune disease; systemic lupus erythematosus; rheumatoid arthritis; multiple sclerosis; haemolytic anaemia; myasthenia gravis; allergic reaction; asthma; inflammatory condition; inflammatory bowel disease; B cell stimulator; T cell activator; blood-related disorder; eosinophilia; thrombosis; thromboembolism; atherosclerosis; myocardial infarction; anglina; anaemia; hyperproliferative disorder; cancer; renal disorder; chronic kidney failure; renal tubular acidosis; kidney stone; cardiovascular disorder; respiratory disorder; human; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design expression and secretion vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunomodulator; immunosuppressive; antiinflammatory; dermatological;
antiarthritic; antirheumatic; neuroprotective; antianaemic; muscular;
antiallergic; antiasthmatic; gastrointestinal; anticoagulant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnostic, forensic, gene therapy and chromosome mapping procedures.
                                                                                                                                                                                                                         08-JUL-1997;
                                                                                                                                                                                                                                                                                        10-OCT-2001; 2001US-00973278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 255 BP; 47 A; 55 C; 53 G; 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 thrombolytic; antiarteriosclerotic; cardiant; cytostatic; nephrotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JUN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            secreted protein polynucleotide seqid 870.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                             97US-0051916P.
97US-0051918P.
97US-0051919P.
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97US-0051928P.
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      -0051929P
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CC expression, in gene therapy or as molecular weight markers. The CC polynucleotides and polypeptides are useful for diagnosing, treating or preventing diseases of the immune system, immunodeficiencies, e.g. CC chediak-Higashi syndrome, autoimmune diseases, e.g. systemic lupus crythematosus, rheumatoid arthritis, multiple sclerosis, haemolytic anaemia or myasthenia gravis, allergic reactions, e.g. asthma, cc inflammatory bowel disease. They can also be used as a stimulator of B cell responsiveness to pathogens or as an activator of T cells. The polynucleotides and polypeptides are also cuseful for treating or preventing blood-related disorders, e.g. confarction, unstable angina or anaemia. They can also be used for treating, preventing or diagnosing hyperproliferative disorders (cancers), renal disorders (chronic kidney failure, renal tubular caidosis or kidney stones), cardiovascular disorders or respiratory
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08-JUL-1997;
08-JUL-1997;
08-JUL-1997;
08-JUL-1997;
                                                                                                                                                                                                                                                           preventing
anemia, inf
kidney fail
                                                                                                                                                                                                                                                                                                                                             Fischer
Lafleur
                                                                                                                                                                            encoding them. The polynucleotides are useful in chromosome identification, for radiation hybrid mapping, in controlling gene identification, for radiation hybrid mapping, in controlling gene
                                                                                                                                                                                                                                                                      New isolated nucleic acid encoding human proteins, upreventing or diagnosing e.g. rheumatoid arthritis, anemia, inflammatory bowel disease, atherosclerosis,
                                                                                                                                                                                                                                                                                                                       WPI; 2004-225733/21.
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                                                                                                                                                                                                              The invention describes novel human secreted proteins and the nucleotides
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ROSEN C A.
SOPPET D R.
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OLSEN H.
EBNER R.
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MOORE P
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ZENG Z.
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Moore
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Shi Y, Olsen
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sen H, Ebner R, Birse
                                                                                                                                                                                                                                                                                   proteins, useful for treating, arthritis, multiple sclerosis,
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Matches 50
 31-JAN-2000

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-JAN-2001; 2001WO-US001354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200157182-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local San
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAGAGACTCTGTCTCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immune/haematopoietic antigen genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immune; haematopoietic; immune/haematopoietic antigen;
atic; gene therapy; vaccine; metastasis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 1.6%;
Similarity 100.0%;
50; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
2000US-01186350P

2000US-018466P

2000US-01846350P

2000US-0198123P

2000US-0199076P

2000US-0199076P

2000US-0199123P

2000US-0214866P

2000US-021664PP

2000US-021664PP

2000US-021649PP

2000US-021649PP

2000US-0217496PP

2000US-0224519P

2000US-0225214PP

2000US-0225214PP

2000US-0225214PP

2000US-0225214PP

2000US-0225214PP

2000US-0225214PP

2000US-0225275PP

2000US-0225275PP

2000US-022575PP

2000US-022575PP

2000US-022575PP

2000US-0225668PP

2000US-0225668PP

2000US-0225668PP

2000US-0225668PP

2000US-0225668PP

2000US-0225668PP

2000US-022575PP

2000US-0225668PP

2000US-022575PP

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 75 C; 80 G; 44 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 50; DB Pred. No. 1.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 12;
1.5e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 288;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ID NO:38904.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ħ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       electronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>0</u>
   01-SEP-2000

01-SEP-2000

01-SEP-2000

01-SEP-2000

06-SEP-2000

08-SEP-2000

08-SEP-2000

08-SEP-2000

08-SEP-2000

14-SEP-2000

14-SEP-2000

14-SEP-2000

14-SEP-2000

14-SEP-2000

14-SEP-2000

12-SEP-2000

14-SEP-2000

12-SEP-2000

14-SEP-2000

12-SEP-2000

12-SEP-2000

13-SEP-2000

21-SEP-2000

22-SEP-2000

22-SEP-2000

23-SEP-2000

24-SEP-2000

25-SEP-2000

20-CCT-2000

20-CCT-2000

20-CCT-2000

20-CCT-2000

20-CCT-2000

20-CCT-2000

20-CCT-2000

02-OCT-2000

03-NOV-2000

04-NOV-2000

08-NOV-2000

08-NOV-2000

08-NOV-2000

08-NOV-2000

08-NOV-2000

08-NOV-2000

01-NOV-2000

2000US-0229343P

2000US-0229345P

2000US-0229345P

2000US-0230437P

2000US-0231244P

2000US-0231244P

2000US-0231244P

2000US-0231241P

2000US-0231411P

2000US-0231411P

2000US-023141P

2000US-023239P

2000US-023239P

2000US-023423P

2000US-0235834P

2000US-0235834P

2000US-0235834P

2000US-0235834P

2000US-0235834P

2000US-0235834P

2000US-0235834P

2000US-0235834P

2000US-024673P

2000US-024673P

2000US-024673P

2000US-024673P

2000US-024652P

2000US-024652P

2000US-024653P

``

2000US-0249213P. 2000US-0249214P. 2000US-0249215P.

```
AAS93725
ID AAS9
XX
AC AAS9
XX
   맑
   Ś
   RESULT 34
  Query Match
Best Local :
  Matches
  17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
01-DEC-2000
01-DEC-2000
05-DEC-2000
05-DEC-2000
06-DEC-2000
06-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
  proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disporders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK84950 and AAM82169 represent sequences used in the exemplification of the present invention
   amino acid sequences given in AAM82170 to AAM91921. (I) have cytos activity, and can be used in gene therapy and vaccine production.
  Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis
                AAS93725;
  Sequence 301 BP;
   Disclosure; SEQ ID NO 38904; 3071pp + Sequence Listing; English.
  AAS93725 standard; cDNA;
   WPI; 2001-483426/52.
  Rosen
   (HUMA-) HUMAN GENOME SCI INC.
  3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  Ç,
   234 AGATTGTGCCACTGCACCTCGGCCAGCAGAGCAGAGCTCTGTCTC
  Similarity
   to AAK64702 encode the human immune/haematopoietic antigen (I) id sequences given in AAM82170 to AAM91921. (I) have cytostatic
  Barash SC,
   2000US-0250160P.

2000US-0250391P.

2000US-0251030P.

2000US-0251988P.

2000US-0256719P.

2000US-02518479P.

2000US-0251868P.

2000US-0251868P.
  larity 100.0%;
Conservative
  2000US-0251990P
2000US-0254097P
   2000US-0249300P.
2000US-0250160P.
   2000US-0249297P.
2000US-0249299P.
  2000US-0251989P
   87
  A; 74 C; 92
   1.6%;
  Ruben SM;
  432 BP
  0
   Score 50;
Pred. No.
   <u>ი</u>
  Mismatches
  48 T; 0 U;
   1.5e-12;
hes 0;
   DB 4;
   0 Other;
   Length 301;
  Indels
  0
  Gaps
  0
```

S 밁

0

04-NOV-2004 ADQ81170; ADQ81170

(first entry)

standard;

DNA; 1001

ВÞ

```
RESULT 35
ADQ81170
ID ADQ81
XX
AC ADQ81
XX
DT 04-NC
XX
   The invention relates to isolated polynucleotide (I) and polypeptide (II) CC sequences. (I) is useful as hybridisation probes, polymerase chain CC reaction (PCR) primers, oligomers, and for chromosome and gene mapping, CC and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed CC in diagnostics as expressed sequence tags for identifying expressed CC genes. (I) is useful in gene therapy techniques to restore normal CC contivity of (II) or to treat disease states involving (II). (II) is contivity of (II) and the bidding partners are useful in medical imaging CC involving aberrant protein expression or biological activity. The CC polypeptide in treating protein expression or biological activity. The CC involving aberrant protein expression or biological activity. The CC involving aberrant protein expression or biological activity. The CC diagnostics, forensics, gene mapping, identification of mutations or responsible for genetic disorders or other traits to assess biodiversity and to produce other types of data and products dependent on DNA and CC amino acid sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention. Note: The sequence data for this CC coding sequences of the invention. Note: The sequence data for this CC electronic format directly from WIPO at CC fit, wipo.int/pub/published_pct_sequences
   Matches
   Query Match
Best Local Similarity
   Sequence 432 BP; 57 A; 142 C; 168 G; 65 T; 0 U; 0 Other;
  New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutation responsible for genetic disorders or other traits and to assess
   WPI; 2001-639362/73.
P-PSDB; ABG29538.
  31-MAR-2000; 2000US-00540217.
23-AUG-2000; 2000US-00649167.
   30-MAR-2001; 2001WO-US008631.
  Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder; ss.
   DNA encoding
  Claim 1; SEQ ID NO 29529; 103pp; English.
  Drmanac RT,
   11-OCT-2001.
   WO200175067-A2.
   Homo sapiens.
   13-FEB-2002
  (HYSE-) HYSEQ INC.
  888 GACGGGCTCAACAAGACGACTGCGTGCTACCACCACCTGGTGCTGACCGT
   50;
   GACGGGCTCAACAAGACGACTGCGTGCTACCACCACCTGGTGCTGACCGT
   Liu C,
   Conservative
  (first entry)
  novel human diagnostic protein #29529
   100.0%;
  Tang
   ĭ
   0,
  Score 50;
Pred. No.
   Mismatches
  DB 5;
  1.5e-12
  Length 432;
   Indels
   937
   0,
   Gaps
```

```
The invention relates to a novel polynucleotide encoded by a phenotype CC associated (PA) gene. The polynucleotide is selected from 292 sequences CC comprising 301-1002 base pairs (AD080913-AD081204) given in the CC specification, with allelic variation contained in a functional CC surrounding like full length cDNA for PA gene polypeptide and with or CC without the PA gene promoter sequence. A polynucleotide of the invention has cardiant activity, and acts as a phenotype-associated gene modulator. The reagent of the invention is useful for preparing a medicament The CC method of the invention is useful for preparing a medicament trailored to suit a patient's individual response to statin therapy. The genetic CC polymorphisms are useful for assessing the response to lipid lowering CC drug therapy and adverse drug reactions of the medicaments, particularly for assessing cardiovascular risks in humans e.g. atherosclerosis, cischaemia/reperfusion, and stroke. The genetic polymorphisms are also useful for identifying compounds for treatments of cardiovascular disease cabove or as prophylactic therapy for cardiovascular diseases. The genetic variations are useful for predicting personal medication schemes omitting adverse drug reactions and allowing an adjustment of the drug dose to cardiove combinant production of normal variant peptides or polypeptides encoded by the genes. The present sequence represents a polynucleotide of the invention.
밁
                                S
   Query Match
Best Local S
Matches 50
   Claim 1;
  cardiovascular diseases.
  New polymorphisms of a phenotype associated (PA) gene, useful for assessing the response to lipid lowering drug therapy and adverse drug reactions of the medicaments, and for screening compounds for treating
   WPI; 2004-581012/56.
   31-JAN-2003; 2003EP-00002212.
03-FEB-2003; 2003EP-00002153.
  atherosclerosis; ischaemia; reperfusion; hypertension; restenosis; arterial inflammation; myocardial infarction; stroke; single nucleotide polymorphism; SNP.
   Sequence 1001 BP; 363
   (FARB ) BAYER HEALTHCARE AG
  23-JAN-2004; 2004WO-EP000539
   WO2004067774-A2
  Homo
   Human phenotype associated polynucleotide baySNP59113
   variation
                                 3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  sapiens
   ٦
  Similarity
   SEQ ID NO 258; 349pp; English.
   phenotype associated; PA; cardiant; statin;
   Schwers S,
   Conservative
   Location/Qualifiers
  /*tag= a
/standard_name= "Single nucleotide polymorphism"
   1.6%;
   A; 150 C;
  Score 50; DB 1; Pred. No. 1.50
   0,
   269 G;
   218 T; 0 U; 1 Other;
   1.5e-12;
hes 0;
   DB 13;
   Length 1001;
   Indels
   cardiovascular;
  SEQ ID NO:258.
   0
   Gaps
   0;
```

RESULT 36

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RESULT 37
ABA82624
ID ABA82
XX
AC ABA82
XZ
AC ABA82
XX
DT 25-JF
XX
XX
Humar
XX
KW Humar
   닭
   ş
   ABS58182
   Query Match
Best Local S
Matches 50
   described is the process for preparing the protein by DNA recombination and the application of the polypeptide and polymucleotide in treating various diseases such as malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases, and various inflammations. The present sequence encodes human zinc finger protein 10.01
  Mao
Human; high bone mass; HBM gene; Zmax1 gene; chromosome 11; 11q13.3; sequence tagged site; STS; osteoporosis; osteopathic; gene therapy;
                                       Human HBM gene region b200e21-h_contig1.
  Sequence 2407
  The present invention relates to the isolation of human zinc fin protein 10.01, and the polynucleotide sequence encoding it. Also
   Claim 6;
  diseases
   New human zinc finger protein 10.01 polypeptide tumors, hemopathy, human immunodeficiency virus
  06-NOV-2000; 2000CN-00127241
   Human; zinc finger protein 10.01; malignant tumour; haemopathy;
human immunodeficiency virus infection; HIV infection; inflammation;
  26-FEB-2003
  ABS58182;
  ABA82624 standard; DNA; 8705 BP
  06-NOV-2000;
   CN1352110-A.
   Homo sapiens.
  immunological
  cDNA encoding human zinc finger protein 10.01
   ABS58182 standard; cDNA; 2407
   3073
   2002-692406/75
DB; ABG72222.
   225
   h 1.6%;
Similarity 100.0%;
   BODE GENE
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGACCAAGACTCTGTCTC
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   Page 25-26 (disclosure); 33pp; Chinese.
  and various inflammations.
  Conservative
  2000CN-00127241
  (first entry)
  (first entry)
   BP;
  disease; gene; ss.
  /product= "Zinc finger protein 10.01"
  Location/Qualifiers
1279. .1554
  /*tag=
   746 A; 458 C; 570 G;
   DEV CO LTD SHANGHAI.
  <u>.</u>
   Score 50;
Pred. No.
  ΒP
  Mismatches
  633
  DB 6;
   1.4e-12;
   T; 0 U; 0 Other;
   Length 2407
   for treating malignant infection, immunological
   3122
   274
  0
  Gaps
```

```
ID XXX XXX PR PR XXX PR
   밁
  ঠ
   Query Match
Best Local S
Matches 50
                         11-MAY-2001;
17-MAY-2001;
01-FEB-2002;
04-MAR-2002;
   Human; high bone mass; HBM; LRP5; LRP6; transgenic; bone mass modulation; gene therapy; bone density modulation; bone strength; trabecular number; bone size; bone tissue connectivity; bone disease; osteoporosis; osteomalacia; rickets; Paget's disease; neoplasm of the bone; gene; ds.
  The present invention describes the human Zmax1 gene and the high bone mass (HBM) gene, which are found on chromosome 11q13.3. The Zmax1 and HBM genes have osteopathic activities. The genes can be used in gene therapy, antisense therapy and in the production of vaccines. They can be used in the diagnosis and treatment of bone disorders including osteoporosis, paget's disease, sclerostosis, osteomalacia and fibrous dysplassia.

ABA82038 to ABA82700 and AAG68168 to AAG68193 represent sequences used in
   05-APR-2000;
05-APR-2000;
  antisense therapy; vaccine; bone disorder; Paget's disease; sclerostosis;
osteomalacia; fibrous dysplasia; ds.
  Carulli
  21-JUN-2000;
   18-OCT-2001.
   WO200292764-A2
  Homo sapiens.
   Human HBM
  ACC45365
  ACC45365 standard; DNA; 8705 BP
   Sequence 8705
   New high bone mass (HBM) modulating bone mass for
  (GENO-)
   6492
  3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  exemplification of the present invention
   1 Similarity 50; Conserv
  GENOME THERAPEUTICS CORP
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   gene fragment #6.
                         ; 2001US-0290071P.
; 2001US-0291311P.
; 2002US-0353058P.
; 2002US-0361293P.
   Conservative (
   2000US-00543771.
2000US-00544398.
  Little
   2000WO-US016951.
  2002WO-US014876.
  (first entry)
  BP;
   303-308; 443pp; English.
  2107
   1.6%; Score 50; DB
100.0%; Pred. No. 1.
ive 0; Mismatches
  A; 2317 C; 2399 G; 1882
  and Zmax1 genes and proteins useful the treatment of e.g. osteoporosis.
  Recker RR,
  Johnson
   1.4e-12;
hes 0;
   몂
   ა
•
  ž
  T; 0
  Length 8705;
  Indels
   U; 0 Other;
  3122
   6541
   <u>,</u>
  for
  Gaps
  0
```

```
CC bone mass (HBM) gene, expressing the corresponding which type HBM gene, comprising an alteration of the gene encoding LRP5 or LRP6, or expressing the target of the transpent of the sequence introduced by homologous or non-homologous recombination. The transgenic animals are for the study of bone density modulation or bone mass modulation. The compressing the invention may have a use in gene therapy. The transgenic animals and composed in more than one parameter selected from bone density, bone species in more than one parameter selected from bone density, bone compositions, which may be employed for treating or preventing bone compositions, which may be employed for treating or preventing bone consplasms of the bone. The transgenic animals and nucleic acids and more compositions, which may be employed for treating or preventing bone compositions, which may be employed for treating or preventing bone compositions of the bone. The transgenic animals and nucleic acids are also compositions of the bone. The transgenic animals and nucleic acids are also compositions of the bone. The transgenic animals and nucleic acids are also compositions of the bone of the invention contains the present sequence is used in the exemplification of the invention.
밁
  δ
   Matches
  Query Match
Best Local
                               11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-0291311P.
01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
   Osteopathic; Gene therapy; High Bone Mass; HBM; LRP5; Zmax1; LRP6; bone mass modulation; osteoporosis; human; ds.
   Sequence 8705 BP; 2107 A; 2317 C; 2399 G; 1882 T; 0 U; 0 Other;
  New transgenic animals (e.g. mice), useful as models for studying bone density modulation, developing drugs for treating or preventing bone diseases (e.g. osteoporosis), or diagnosing diseases characterized by
   04-DEC-2003
  Example 2; Page 358-361; 603pp; English.
   ADB98065
   13-MAY-2002; 2002WO-US014877
  21-NOV-2002.
   WO200292000-A2
  HBM-related clone contig b200e21-h contig1
  (GENO-) GENOME THERAPEUTICS (AMHP) WYETH.
   invention
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   50;
   bone
  Similarity
   standard;
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   Bex FJ,
   1.6%; Sc
llarity 100.0%; F
Conservative 0;
   density.
   (first entry)
   relates to novel transgenic animals expressing the y gene, expressing the corresponding wild type HBM
   DNA;
   Yaworsky PJ,
  CORP
  Score 50;
Pred. No.
   Mismatches
  Bodine PV;
  B
  1.4e-12;
  8
  Length 8705;
   Indels
   6541
  3122
   0
  bone
d by
   Gaps
```

(GENO-)

GENOME THERAPEUTICS CORP

```
RESULT 40
  밁
   Ś
   Query Match
Best Local S
Matches 50
  Allen K,
Yaworsky
                                  Regulating LRP5, LRP6 or HBM activity in a subject, useful for modulating lipid levels and/or bone mass, and for in treating bone mass disorders, e.g. osteoporosis, comprises administering a composition which modulates
   17-MAY-2001; 2001US-0291311P.
01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
   17-MAY-2002; 2002WO-US015982.
  21-NOV-2002
   WO200292015-A2
   Homo sapiens
  hyperostosis;
   29-JAN-2004
   ADE82434;
  ADE82434 standard;
  Sequence 8705
  phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence was used to illustrate the invention.
   suffering
   New nucleic acid comprising a mutation in LRP5 or LRP6, useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject
  WPI; 2003-129214/12.
   WPI; 2003-129219/12
   Antiarthritic;
  LRP5; LRP6; HBM; Dkk activity; Osteopathic; Antiinflammatory;
   Example 3; SEQ ID NO 10; 629pp; English.
  (AMHP ) WYETH
  (GENO-)
   6492
   3073
                            activity.
  DNA sequence
  ۲
   GENOME WYETH.
  Similarity
  Anisowicz
PJ;
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  from e.g. osteoporosis.
   Anisowicz
   Conservative
  (first entry)
   BP;
   osteogenesis;
  THERAPEUTICS
   bone mass disorders;
  2107 A; 2317 C;
  related to
   DNA;
   þ
  ۶
   1.6%;
   Bhat
  8705
  Graham
  CORP
   Wnt
   0
   Score :
   8
   BP
  the
  됬
   вignaling;
   Mismatches
   2399
   Damagnez
  invention
   . No.
   osteoporosis; hypercalcaemia;
   Morales
   G; 1882
   1.4e-12;
  DB 10;
   ۲,
   da.
  Þ
   0
   Robinson
   T; 0 U; 0 Other;
   Length 8705;
  Yaworsky
   Indels
  ď,
  6541
  <u>.</u>
  Liu
   Gaps
  Σ
  0
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Disclosure;

SEQ

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10;

173pp;

English

The invention relates to an isolated amino acid protein sequence selected from an amino acid sequence appearing as ADR16922 or an amino acid sequence comprising or consisting of the extracellular domain of ADR16922 (amino acids 23-1385). ADR16922 is encoded by the HBM (high bone mass) allele of the human zmax1 gene and has sequence similarity to LDL receptors. Also disclosed are nucleic acids, proteins, cloning vectors, expression vectors, transformed hosts, methods of developing pharmaceutical compositions, methods of identifying molecules involved in bone development, and methods of diagnosing and treating diseases involved in bone development. Specifically disclosed is the Zmax1 gene

Example 2; SEQ ID NO 10; 284pp; English.

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밁
   δ
  Matches
  Best
  Query Match
  13-JAN-1998;
23-OCT-1998;
13-JAN-1999;
  The present invention relates to regulating LRP5, LRP6 or HBM activity is a subject comprising administering a composition which modulates a Dkk activity. The method is useful for modulating lipid levels and/or bone mass, and is useful in treating or diagnosing abnormal lipid levels and bone mass disorders, such as osteoporosis, bond fracture, age-related loss of bone, a chondrodystrophy, drug-induced bone disorder, high bone turnover, hypercalcaemia, hyperostosis, osteogenesis, imperfecta, osteomalacia, osteomyelitis, Paget's disease, osteoarthritis, and rickets. Modulators of Dkk activity are useful for as reagents in studying bone mass and lipid level modulation, in modulating Wnt signaling, or treating Dkk-mediated disorders. The present sequence represents a human DNA sequence related to the invention.
   New high bone mass gene of chromosome 1.1Q13.3, encoding protein useful for treating, diagnosing, preventing, or screening for normal and abnormal conditions of bone, including metabolic bone diseases, e.g.
  chromosome 11q13.3; osteopathic; LDL receptor; bone devemetabolic bone disease; bacterial artificial chromosome.
  WPI; 2004-623529/60.
  Carulli JP,
  05-APR-2000;
  US6780609-B1.
  Human; high bone mass; Zmax1; ds; BAC; HBM; osteoporosis; chromosome 11q13.3; osteopathic; LDL receptor; bone devel
  BAC clone containing
  04-NOV-2004
  ADR16928
  Sequence
   24-AUG-2004.
   ADR16928;
  (GENO-) GENOME THERAPEUTICS CORP.
   Local Similarity
  3073
  6492 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   sapiens.
  50;
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  8705 BP;
  standard; DNA; 8705
  Conservative
   Little
  2000US-00543771.
   (first
  98US-0071449P.
98US-0105511P.
99US-00229319.
  2107 A; 2317 C; 2399 G; 1882 T; 0 U; 0 Other;
   ₽,
   entry)
  segments
  100.0%;
   1.6%;
   Recker
  0
   Score 50;
Pred. No.
   ΒP
  of the human
   ₽
,
  Mismatches
   Johnson
  1.4e-12;
   DB 10;
   Zmax1
   ž
   bone development;
  Length 8705;
  gene
  Indels
   #6
  6541
  3122
   0;
   Gaps
  ä
   0
```

```
ARSULT 42
ADR47579
ID ADR47579
AC ADR47
  នននិន្នន
   밁
   S
CC high bone mass protein (HBM). The gene exists in two alleles, Zmaxi, the contional wild-type (the cDNA for which appears as ADR47570 encoding ADR47572) and the HBM allele (the cDNA for which appears as ADR47571 encoding ADR47573). The two alleles differ by a single nucleotide conoding ADR475731. The two alleles differ by a single nucleotide change at position 171 of the protein. Also included are a replicative clange at position 171 of the protein. Also included are a replicative clange at position in a protein (and a replicative in an isolated host cell), an expression vector comprising HBM/Zmaxi operably linked to a transcription regulatory region, an isolated host cell transformed with the vector(s), a method for testing a substance as a charageutic agent for bone modulation in a host, a method of identifying a claudidate) protein involved in bone modulation, a method of testing for C (candidate) protein involved in bone modulation, a method of testing for HBM activity, a method of development disorders, a method for treating a bone development disorders, a method of altering bone development in a host, a
   Query Match
Best Local S
Matches 50
  and the high bone mass (HBM) allele on chromosome 11q13.3 encoding ADR16922. The protein is useful for treating, diagnosing, preventing, or screening for normal and abnormal conditions of bone, including metabolic bone diseases, e.g. osteoporosis. The present sequence is a BAC (bacterial artificial chromosome) containing part of the Zmax1 gene.
  Sequence 8705 BP; 2107 A; 2317 C; 2399
   13-JAN-1999; 99US-00229319
05-APR-2000; 2000US-00544398
  13-JAN-1998;
23-OCT-1998;
  09-SEP-2004.
  New nucleic acid sequence encoding high
  Carulli JP,
   10-DEC-2003;
   US2004176582-A1
   bone modulation; bone development disorder; osteoporosis; chromosome 11q13.3; gene therapy; BAC.
  BAC clone
   02-DEC-2004
   ADR47579 standard; DNA; 8705 BP
  invention relates to
   6492 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  2004-661408/64
   sapiens.
  50;
  ds; bacterial artificial chromosome; high bone mass;
  GENOME THERAPEUTICS UNIV CREIGHTON.
   Similarity
  and/or preventing osteoporosis
  containing segments of the human
  SEQ ID NO 10; 303pp; English.
  Conservative
  Little RD,
   2003US-00731739
   (first entry)
  98US-0071449P.
98US-0105511P.
99US-00229319.
  1.6%;
  an isolated nucleic acid sequence encoding a
   Recker RR,
  0
   CORP
   Score 50;
Pred. No.
  Mismatches
  Johnson
  G; 1882 T; 0
   DB
   bone
  1.4e-12;
  Zmax1 gene #6.
   mass,
   ĭ
  ç
   Length 8705;
  Indels
   useful in diagnosing,
  U; 0 Other;
   6541
   Zmax1;
  Gaps
  0
```

```
RRESULT 43
AREB6930AB
AID 93BB69
AX
AEB693CA
AX
AEB693CA
AX
AEB693CA
AX
AEB693CA
AX
AEB693CA
AX
AEB66
AX
AEB66
AX
AEB66
AX
AEB66
AEB66
AEB693CA
AEB66
AEB693CA
AEB66
AEB693CA
AEB66

   맑
  5
  Query Match
Best Local S
Matches 50
   method for diagnostic screening for a genetic predisposition to a bone development disorder, a diagnostic assay for bone development disorders, a method of expressing the HBM protein in bone tissue, a bacterial artificial chromosome comprising HBM//max1 sequence (appearing as an DR47574-ADR47580), a method for amplifying a nucleotide polymorphism in the Zmax1 or HBM gene, a method for identifying a regulatory element of a HBM gene and an isolated nucleic acid segment of at least 15 contiguous nucleotides including a polymorphic site from HBM/Zmax1. The nucleic acid molecule and the encoded polymorphic site from HBM/Zmax1. The nucleic acid in diagnosing, treating and preventing a bone development disorder, i.e. osteoporosis. The gene for HBM/Zmax1 is located on chromosome 11q13.3. The present sequence is an HBM DNA from a bacterial artificial
  13-JAN-1998;
23-OCT-1998;
13-JAN-1999;
                        candidate molecule involved in bone modulation. The method comprises identifying a molecule that binds to High Bone Mass protein (HBM) and/Zmax1 protein. The HBM gene exists in two alleles: Zmax1, the notional wild-type (the cDNA for which appears as AEB69299 encoding AEB69301 and the HBM allele (the cDNA for which appears as AEB69300 encoding AEB69300). The two alleles differ by a single nucleotide polymorphism (T to G at position 582 of AEB69299)
  Sequence 8705
   Osteopathic; high bone mass; Zmax1; bone disease; osteoporosis;
   AEB69308
  chromosome.
  US2005142617-A1
   Homo sapiens.
   osteomalacia;
   Human High Bone Mass gene related contig b200e21-h_contig1,
   22-SEP-2005
  The present invention relates to a method (M1) for identifying a
   Example
   WPI; 2005-496364/50.
  Carulli JP,
  29-APR-2004; 2004US-00834377
  Identifying candidate molecule involved in bone modulation, identifying molecule that binds to Zmax1, high bone mass (H
   (UYCR-)
  3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGAGCTCTGTCTC 3122
   Similarity
   VIND
   2; SEQ ID NO 10; 308pp;
   Zmax1 and HBM
  GENOME THERAPEUTICS CORP
   standard;
   AGATTGTGCCACTGCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 6541
   1.6%;
ilarity 100.0%;
Conservative
  Little RD,
   2000US-00543771.
  CREIGHTON SCHOOL MEDICINE
  (first entry)
  BP; 2107 A; 2317 C; 2399 G; 1882
   bone
  98US-0105511P.
99US-00229319.
  98US-0071449P
   DNA;
   injury; Pagets disease;
   change at position 171
   protein.
  8705
  Recker RR, Johnson ML;
  0;
   Score 50; pred. No.
   English.
   Mismatches
  1.4e-12;
  high bone mass (HBM)
  13;
  T; 0 U; 0 Other;
  Length 8705;
  Indels
  SEQ
   0
  protein,
  Ħ
  and
   ů,
  0
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causing a Gly

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RESULT 44
AAK866119
ID AAK866
XX AAK86
XX AAK86
XX Humar
XW Humar
XW Cytos
XX Homo
XX Homo
XX Homo
XX Homo
XX Homo
XX Homo
XX 17-JA
PR 11-JH
PR 11-
   문
   δ
   នន្តន្តន្តន្តន្ត្
  Matches
   Query Match
  31-JAN-2000;
04-FBB-2000;
04-FBB-2000;
02-MAR-2000;
11-MAR-2000;
11-MAR-2000;
11-MAR-2000;
07-JUN-2000;
07-JUL-2000;
07-JU
  protein has the property of causing elevated bone mass, while the Zmaxl protein does not. The gene for HBM/Zmaxl is located on chromosome 11q13.3. Also claimed is a method of pharmaceutical development for treating of bone development disorders, such as osteoporosis, osteomalacia, bone fractures, Paget's disease, etc., which comprises identifying a molecule that binds to the Zmaxl protein, or to HBM, or both. The present sequence was used to illustrate the invention.
  17-JAN-2001;
   09-AUG-2001.
  WO200157182-A2
  cytostatic;
  Human;
   Human immune/haematopoietic antigen
  07-NOV-2001
   AAK86119;
  Sequence 8705 BP; 2107 A; 2317 C; 2399
  AAK86119
  Local Similarity
   50;
  immune; haematopoietic; immune/haematopoietic
  standard;
2000US-0179065P

2000US-0184664P

2000US-0184664P

2000US-0184664P

2000US-0198076P

2000US-0199076P

2000US-0199076P

2000US-0214886P

2000US-0216840P

2000US-0216840P

2000US-0217487P

2000US-0217487P

2000US-021889P

2000US-021829P

2000US-021829P

2000US-0221829P

2000US-0224518P

2000US-0224518P

2000US-0225713P

2000US-0225266P

2000US-0225267P

2000US-02252447P

2000US-0225758P

   gene
  1.6%; Score 50; DB ilarity 100.0%; Pred. No. 1.. Conservative 0; Mismatches
  2001WO-US001354.
   (first
  therapy; vaccine; metastasis;
   DNA;
   entry)
  10396
  BP.
   genomic sequence
   G; 1882 T; 0 U; 0 Other;
   DB 14;
1.4e-12
   .4e-12;
s 0;
   gb
  Length 8705;
  Indels
   antigen; cancer;
  SEQ ID NO: 40931.
   ç,
   Gaps
  0
       $\,\text{$\frac{1}{2}\text{$\frac{1}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}{2}\text{$\frac{1}\text{$\frac{1}\text{$\frac{1}\text{$\frac{1}\text{$\frac{1}\text{$\frac{1}\text{$\frac
  22-AUG-2000
23-AUG-2000
30-AUG-2000
01-SEP-2000
01-SEP-2000
05-SEP-2000
06-SEP-2000
06-SEP-2000
08-SEP-2000
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14-SEP-2000
15-SEP-2000
16-NOV-2000
17-NOV-2000
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10-NOV-2000
10-NOV-2000
10-NOV-2000
10-NOV-2000
10-NOV-2000
10-NOV-2000
11-NOV-2000
2000US-0227182P.
2000US-0228924P.
2000US-0229343P.
2000US-0229343P.
2000US-0229343P.
2000US-0229343P.
2000US-0231043P.
2000US-0231244P.
2000US-0231244P.
2000US-0231413P.
2000US-0231414P.
2000US-023144P.
2000US-023144P.
2000US-023144P.
2000US-023144P.
2000US-023144P.
2000US-023144P.
2000US-023144P.
2000US-0231499P.
2000US-0241182P.
2000US-024647P.
2000US-024647P.
2000US-024647P.
2000US-024652P.
2000US-024652P.
2000US-024652P.
2000US-024651P.
2000US-024661P.
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                        Ś
  Matches
  Query Match
Best Local Similarity
  17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
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17-NOV-2000
17-NOV-2000
01-DEC-2000
01-DEC-2000
05-DEC-2000
05-DEC-2000
06-DEC-2000
08-DEC-2000
  AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAW62170 to AAW91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disporders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-crived cells. AAK64703
  Sequence 10396
  sequences
  6
  Disclosure; SEQ ID NO 40931; 3071pp + Sequence Listing; English
   Nucleic
   Rosen
  AAK87694 represent human immune/haematopoietic antigen genomic quences from the present invention. AAK54942 to AAK54950 and AAM82169
 5597
                           3073
   2001-483426/52.
   ÇĄ,
   for
   acids encoding
  HUMAN
               AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 5646
  sequences
  preventing,
   Barash
  2000US-0249211P.
2000US-0249213P.
2000US-0249214P.
2000US-0249215P.
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2000US-0249216P.
2000US-0249216P.
2000US-0249216P.
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2000US-0249264P.
2000US-0249264P.
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2000US-0249299P.
2000US-025936P.
2000US-025936P.
2000US-025936P.
2000US-0251866P.
2000US-0251866P.
  2000US-0249209P.
2000US-0249210P.
2000US-0249211P.
   2000US-0251989P.
2000US-0251990P.
2000US-0254097P.
  Conservative
  2001US-0259678P
  GENOME
  BP; 3175 A;
   SC,
  used in the exemplification
  1.6%;
  SCI
  human immune/hematopoietic antigen polypeptides, diagnosing and/or treating cancers and metastasis.
   Ruben SM;
  2158 C;
  0
  Score 50;
Pred. No.
  Mismatches
  2127
  Ģ.
  멂
   1.4e-12;
  4.
  2936 T; 0 U; 0 Other;
  0
   Length 10396;
   of the
  Indels
  present invention
   0
   Gaps
   0
```

RESULT 45 ABA20857

```
Human nervous system related polynucleotide SEQ ID
   23-JAN-2002
   ABA20857 standard; DNA; 11234
   (first entry)
   BP.
  NO 13188
```

Human; nootropic; neuroprotective; cytostatic; dermatological; viri immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulnu antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer, antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder; neurological disease; infection; nephrotropic; gene therapy; vaccine; antiarthritic; cancer; dermatological; virucide; antibacterial; vulnerary; ds.

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17.MAR-2000
19.MAY-2000
19.MAY-2000
28.JUN-2000
27.JUN-2000
30.JUN-2000
07.JUL-2000
11.JUL-2000
11.JUL
   24-FEB-2000;
02-MAR-2000;
  31-JAN-2000;
   16-AUG-2001.
   17-JAN-2001;
   WO200159063-A2
2000US-0179065P.
2000US-0184664P.
2000US-0184664P.
2000US-0184664P.
2000US-0199076P.
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2000US-0214886P.
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2000US-0216489P.
2000US-0216489P.
2000US-0217487P.
2000US-0224518P.
2000US-0224519P.
2000US-0225214P.
2000US-0225214P.
2000US-0225214P.
2000US-0225266P.
2000US-022526P.
2000US-0225759P.
2000US-0235759P.
2000US-
   2001WO-US001334
```

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   08-SEP-2000
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11-SEP-2000
11-SEP-2000
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29-SEP-2000
29-SEP-2000
20-OCT-2000
02-OCT-2000
01-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
01-NOV-2000
2000US-0236370P

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2000US-0237033P

2000US-0237039P

2000US-0237039P

2000US-0237039P

2000US-0241786P

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2000US-0241808P

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2000US-0246477P

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2000US-0246478P

2000US-0246478P

2000US-0246617P

2000US-0246611P

2000US-0246528P

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2000US-0249214P

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2000US-0249214P

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2000US-0249245P

2000US-0249299P

2000US-0249299P
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2000US-0232081P.
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2000US-023196P.
2000US-0232397P.
2000US-023239P.
2000US-023239P.
2000US-023239P.
```

RESULT 46 AAKB0184/c ID AAK80184 £

standard; DNA; 13026

BP

0,

cytostatic; gene therapy; vaccine; metastasis;

Human; immune; haematopoietic; immune/haematopoietic antigen; cancer Human immune/haematopoietic antigen genomic sequence SEQ ID NO:34996

WO200157182-A2 Homo sapiens. 07-NOV-2001 AAK80184;

(first entry)

```
밁
              á
  Query Match 1.6%; Sometive 1.00.0%; Figure 1.00.0%; Particles 50; Conservative 0;
  17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
08-DEC-2000;
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08-DEC-2000;
08-DEC-2000;
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11-DEC-2000;
   Sequence 11234 BP; 3094 A; 2417 C; 2869 G; 2854 T; 0 U; 0 Other;
  Disclosure; SEQ ID NO 13188; 1701pp + Sequence Listing; English
  from WIPO at
  useful for
  Nucleic acids encoding
   (HUMA-)
11155
  metastases.
               3073
  2001-541565/60.
   ÇĄ,
   HUMAN
        AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGACAAGACTCTGTCTC 3122
AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 11204
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2000US-0251869P.
2000US-0251869P.
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2000US-0251869P.
2000US-0251869P.
  ftp.wipo.int/pub/published_pct_sequences
   2001US-0259678P
  GENOME
   SCI
   3224 human nervous system antigen polypeptides, diagnosing and/or treating nervous system cance
   Ruben SM
   INC
                               Score 50; DB;; Pred. No. 1.4:0; Mismatches
  DB 5; Le
                                 <u>,</u>
  Length 11234;
                                 Indels
                                0;
                                Gaps
  cancers
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| PR<br>PR                                                 | 2                                                        | 8 8 8                                | 1 P X                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | PR                                   | PR               | <b>R</b> R                           | PR               | אם<br>אם         | PR               | PR               | אָק              | PR               | PR :             | ק<br>ק           | 2 2              | 2 2              | 뮸                | PR               | PR               | PR               | р;              | 7 G              | 2 2              | PR               | PR               | PR               | PR               | ב<br>ב<br>ב      | ב<br>ב<br>ב      | ž                | PR               | PR               | PR               | PR               | P ;              | ָ<br>ק            | PR               | PR               | PR               | PR ?            | 8 7              | 8 3                                        | 3 5              | PR               | PR               | PR               | PR               | PR               | P. 7             | g ;              | PR        | PR               | XX               | ž        | PD           |
|----------------------------------------------------------|----------------------------------------------------------|--------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------|------------------|--------------------------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|-----------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|-------------------|------------------|------------------|------------------|-----------------|------------------|--------------------------------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|------------------|-----------|------------------|------------------|----------|--------------|
| 2000                                                     | 7-SEP-2                                                  | 5-SEP-2000                           | 1-SEP-2000                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | 4-SEP-2                              | 4-SEP-2000       | 4-SEP-2                              |                  | 2-SEP-2          | 8-SEP-2          |                  | 8-SEP-2          | 8-SEP-2000       | 8-SEP-2          | 8-SEP-2000       | 0000-488-9       | 5-SEP-2000       | 5-SEP-2          |                  |                  |                  |                 |                  |                  |                  |                  |                  |                  | ώŇ               | ÖK               | s ĸ              | 3 N3             | ல்               | Ň                | AUG-2            | 14-AUG-2000:     |                   | 10L-2            | JUL-200          | 200              | N               | 38               | 7-701-200                                  | 0-JUN-200        | 2                | -200             | 9-MAY-200        | 200              | 200              | 200              | 9 6              | 4-FEB     | 200              | 17-JAN-2001;     | !        | 09-AUG-2001. |
| 2000US-0235836P.<br>2000US-0236327P.<br>2000US-0236367P. | 2000US-0235484P.<br>2000US-0235484P.<br>2000US-0235834P. | 2000US-0234274P.<br>2000US-0234997P. | 2000US-0233065P.<br>2000US-0234223P.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | 2000US-0233063P.<br>2000US-0233064P. | 2000US-0232401P. | 2000US-0232399P.<br>2000US-0232400P. | 2000US-0232398P. | 2000US-0231968P. | 2000US-0232081P. | 2000US-0232080P. | 2000US-0231413P. | 2000US-0231244P. | 2000US-0231243P. | 2000US-0230438F. | 2000US-0230437P. | 2000US-0229513P. | 2000US-0229509P. | 2000US-0229345P. | 2000US-0229344P. | 2000US-0229343P. | 2000US-0229287P | 2000US-0227UUSF. | 2000US-0227182F. | 2000US-0226868P. | 2000US-0226681P. | 2000US-0226279P. | 2000US-0225759P. | 2000US-0225758P. | 200005-02254476. | 2000US-02254/UP. | 2000US-0225268P. | 2000US-0225267P. | 2000US-0225266P. | 2000US-0225214P. | 2000US-0225213P. | 2000000-02245160. | 2000US-0220964P. | 2000US-0220963P. | 2000US-0218290P. | 2000US-0217496P | 200005-0216660F. | 2000US-021664/F.                           | 2000US-0215135P. | 2000US-0214886P. | 2000US-0209467P. | 2000US-0205515P. | 2000US-0198123P. | 2000US-0190076P. | 2000US-0189874P. | 200000-0101001F. | 200       | 2000US-0179065P. | 2001WO-US001354. |          |              |
|                                                          |                                                          |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |                                      |                  |                                      |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                 |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                   |                  |                  |                  |                 |                  |                                            |                  |                  |                  |                  |                  |                  |                  |                  |           |                  |                  |          |              |
|                                                          |                                                          |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |                                      |                  |                                      |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                 |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                   |                  |                  |                  |                 |                  |                                            |                  |                  |                  |                  |                  |                  |                  |                  |           |                  |                  |          |              |
|                                                          |                                                          |                                      |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |                                      |                  |                                      |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  |                  | -                |                 |                  |                  |                  |                  |                  |                  | _                |                  |                  |                  |                  |                  |                  |                  |                   |                  |                  |                  |                 |                  |                                            |                  |                  |                  |                  |                  |                  |                  |                  |           |                  |                  |          |              |
| DR XX                                                    | X Z X                                                    | 2 2 2                                | 2 8                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           | PR                                   | PR               | g g                                  | PR :             | 멅                | PR               | PR               | P 20             | PR               | PR ;             | <b>2</b> 5       | 3 5              | 3 2              | 뀵                | PR               | PR               | 뀱 :              | 닭 ;             | 8 8              | 2 2              | R                | PR               | PR               | 뀵                | 2 7              | 2 3              | 3 5              | 2 2              | R                | Ŗ                | PR               | æ ;              | 8 3               | 꿇                | PR               | 멅                | ¥ ;             | מי<br>ק          | ֓֞֝֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֓֡֓֓֡֓֓֡֓֓֓֓֓֡֓֓֡ | 2                | PR               | PR               | PR               | PR               | БК               | 2 3              | ָרָ ק            | 3 3       | PR               | <b>8</b> %       | 뮸        | PR           |
| Rosen CA, B<br>WPI; 2001-48                              | (HUMA-) HUMAN                                            | 1-200                                | 0 HG - 8 - 1 HG - 1 HG - 8 - 1 HG - 1 HG - 8 - 1 HG - 1 | -200                                 | 8-DEC-200        | 6-DEC-200                            | 5-DEC-200        | 1-DEC-200        | 1-DEC-200        | 7-NOV-200        | 7-200            | 7-200            | 7-200            | 7-200            | 200              | 7-200            | 7-200            | 7-200            | 7-200            | 7-200            | 7-200           | 17-NOV-2000;     | 200              | 200              | 200              | 8-NOV-200        | 8-NOV-200        | 8-NOV-200        | 200              | 002-VON-8        | 8-NOV-200        | 8-NOV-200        | 8-NOV-200        | 8-NOV-200        | 200              | 8-NOV-200         | 8-NOV-200        | 8-NOV-200        | 8-NOV-200        | 200             | 0-001-200        | 0-001-200                                  | 0-OCT-200        | 0-OCT-200        | 0-OCT-200        | -200             | 0-OCT-200        | 200              | 3-0CT-200        |                  | 2-OCT-200 | 200              | 200              | -SEP-200 | -SEP-200     |
| Barash SC, Ruben SM;<br>483426/52.                       | N GENOME SCI INC                                         | SOTOO                                | 2000US-0251990P.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | SOOOO                                | snooo            | 80000<br>80000                       | S0000            |                  | snooo            | S00008           | SOCOO            | snooo            | Snooo            | 20000            | SOOO             | SOCOOS           | Snooo            | snooo            | Succo            | 8000s            | S0000           |                  | SOCIA            | SOCO             | Snooo            | SOOOO            | Succe            | Snooo            | Social           | SOUG             | SDOODS           | Snooo            | SOOOO            | SOOOO            | 20000            | 2000              | spood            | S0000            | suooo            | Snood           |                  |                                            | SOCOO            | Sucoo            | Snooo            | Snooo            | SUOOO            | SDOOO            | Snooo            |                  | SDOOO     | 80000            | S0000            | Snooo    | suooo        |

XFFX8X5555555555555555555555

```
RESULT 47
AAX80185/c
ID AAX801:
XX AAX801:
XX AAX801:
XX Human:
XX
  CC amino acid sequences given in AAM82170 to AAM91921. (1) have cytostatic cartivity, and can be used in gene therapy and vaccine production. (1) CC activity, and can be used in gene therapy and vaccine production. (1) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (1) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (1) by expressing inactive proteins or to Supplement the patients own production of (1). Additionally, (1) CC supplement the patients own production of (1). Additionally, (1) CC concers and treat immune/haematopoietic sell to express the CC protein. (1) proteins and polynucleotides may be used to prevent, CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic antigen genomic CC sequences from the present invention. AAK54942 to AAK54950 and AAM62169 CC represent sequences used in the exemplification of the present invention.
   Matches 50;
   Query Match
Best Local
   31-JAN-2000
04-FEB-2000
24-FEB-2000
02-MAR-2000
16-MAR-2000
17-MAR-2000
18-APR-2000
19-MAY-2000
07-JUN-2000
07-JUN-2000
07-JUN-2000
07-JUN-2000
07-JUN-2000
11-JUN-2000
07-JUN-2000
   09-AUG-2001
   WO200157182-A2
   Homo sapiens.
   cytostatic; gene therapy; vaccine; metastasis; ds.
  Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
   Human immune/haematopoietic antigen genomic sequence SEQ ID NO:34997.
   07-NOV-2001
   AAK80185;
   AAK80185 standard; DNA; 13026 BP
   Sequence 13026 BP; 4098 A; 2489 C; 2384 G; 4055 T; 0 U; 0 Other;
  Disclosure; SEQ ID NO 34996; 3071pp + Sequence Listing; English.
  Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis.
  17-JAN-2001;
   Local Similarity
   8926
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCCAAGAGCTCTGTCTC 3122
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGAGCTCTGTCTC 8877
; 2000US-0179065P.

2000US-0184664P.

2000US-0184635P.

2000US-0186350P.

2000US-0199076P.

2000US-0199076P.

2000US-0199076P.

2000US-029467P.

2000US-021647P.

2000US-021647P.

2000US-021647P.

2000US-0217496P.

2000US-0217496P.

2000US-0217496P.

2000US-0218290P.

2000US-0218290P.

2000US-0218290P.
  ilarity 100.0%;
Conservative (
  2001WO-US001354.
  (first entry)
   1.6%;
  0; Mismatches
   Score 50;
Pred. No.
   DB 4; Lo
   0,
  Length 13026;
   Indels
  0;
  Gaps
   0
          \(\text{\text{$\
      114-SEP-2000;
21-SEP-2000;
21-SEP-2000;
25-SEP-2000;
26-SEP-2000;
27-SEP-2000;
29-SEP-2000;
29-OCT-2000;
20-OCT-2000;
21-OCT-2000;
20-OCT-2000;
20-O
   14-AUG-2000;
18-AUG-2000;
22-AUG-2000;
22-AUG-2000;
23-AUG-2000;
30-AUG-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
01-SEP-2000;
05-SEP-2000;
06-SEP-2000;
06-SEP-2000;
08-SEP-2000;
08-SEP-2000;
08-SEP-2000;
08-SEP-2000;
   08-SEP-2000;
08-SEP-2000;
12-SEP-2000;
  14-SEP-2000;
14-SEP-2000;
14-SEP-2000;
14-SEP-2000;
   14-SEP-2000
14-SEP-2000
   14-AUG-2000;
14-AUG-2000;
  14-AUG-2000;
14-AUG-2000;
  14-AUG-2000;
14-AUG-2000;
      2000US-022964P.
2000US-0224519P.
2000US-0224519P.
2000US-0225219P.
2000US-0225266P.
2000US-0225270P.
2000US-0225757P.
2000US-0225759P.
2000US-0225759P.
2000US-0225759P.
2000US-0225759P.
2000US-0225759P.
2000US-0225759P.
2000US-0225759P.
2000US-0229349P.
2000US-0229349P.
2000US-0229343P.
2000US-0231243P.
2000US-023124180P.
2000US-0244474P.
2000US-02446474P.
2000US-02446474P.
```

밁

```
AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) canning acid sequences given in AAM62170 to AAM91221. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, cancers and cancer metastases of haematopoietic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic
  08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
17-NOV-2000
17-NOV
   Nucleic
  Disclosure;
  Rosen
   (HUMA-)
  Ş
   for preventing,
  acids encoding
   HUMAN
  Barash
  2000US-0249219P
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249215P
2000US-0249215P
2000US-0249215P
2000US-0249215P
2000US-0249216P
2000US-0249244P
2000US-0249244P
2000US-0249264P
2000US-0249265P
2000US-0249265P
2000US-0249269P
2000US-0249299P
2000US-0249299P
2000US-0251989P
2000US-0251869P
2000US-0251869P
2000US-0251989P
2000US-0251989P
2000US-0251989P
2000US-025199PP
2000US-025199PP
2000US-025199PP
2000US-025199PP
2000US-025199PP
   2000US - 0246476P.
2000US - 0246477P.
2000US - 0246478P.
2000US - 0246523P.
2000US - 0246524P.
2000US - 0246525P.
2000US - 0246527P.
2000US - 0246528P.
  SEQ ID NO
   GENOME
  SC,
   SCI
  34997; 3071pp + Sequence Listing;
  human immune/hematopoietic antigen polypeptides, diagnosing and/or treating cancers and metastasis.
   INC
  S
```

English.

```
RASULT 48
AALO544
AC AALO54
AC CANCER
CO O2 AUG
CO O2 AUG
CO O2 AUG
CO O3 O4 PEBB
CO O4
  밁
   ঠ
  ឧឧ
   Matches
   Query Match
16-MAR-2000
18-APR-2000
18-APR-2000
19-MAY-2000
07-JUN-2000
07-JUL-2000
07-JUL-2000
11-JUL-2000
11-JUL-2000
11-JUL-2000
14-JUL-2000
14-AUG-2000
15-AUG-2000
16-AUG-2000
17-AUG-2000
18-AUG-2000
18-AUG-2000
18-AUG-2000
18-AUG-2000
19-AUG-2000
  Human reproductive system related antigen DNA SEQ ID NO:
   21-NOV-2001 (first
  AAL05461;
   AAL05461
  Sequence 13026 BP;
   sequences from the present invention. AAK54942 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention
   17-JAN-2001;
   WO200155320-A2
   Homo sapiens.
   cancer;
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   8926 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   reproductive system related antigen; reproductive system disorder;
  50;
   gene therapy;
   Similarity
   standard;
   2000US-0189874P

2000US-019812BP

2000US-0205515P

2000US-0214886PP

2000US-0214887P

2000US-0216847P

2000US-0216847P

2000US-021899P

2000US-021899P

2000US-021899P

2000US-0228519P

2000US-0224519P

2000US-0224519P

2000US-0225213P

2000US-0225759P

2000US-0226868P

2000US-0226868P
  Conservative
2000US-0227009P.
2000US-0228924P.
2000US-0229287P.
2000US-0229343P.
  2001WO-US001339
   DNA;
  4098 A; 2489 C;
   entry)
   1.6%;
  31474
  0,
  Score 50;
Pred. No.
   ВP
  Mismatches
  2384
  DB 4; Le
1.4e-12;
  Ģ
  4055 T; 0 U; 0 Other;
  ç,
   Length 13026;
  Indels
   8149
   8877
  3122
  0;
  Gaps
```

```
RESULT 49
ABL98314/c
ID ABL983
XX ABL983
XX ABL983
XX ABL983
XX ABL983
XX Human
XX Human;
XW reprod
XW reprod
XW cardio
XW gastro
XX Homo 8
XX Homo 8
XX PN WO2001
   밁
  8
   Query Match
Best Local S
Matches 50
  17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
01-DEC-2000
01-DEC-2000
05-DEC-2000
   Isolated used in p
  Human; testicular antigen; testes; cancer; metastasis; immune disorder; reproductive system disorder; urinary system disorder; gene therapy; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disease; infection; cytostatic; gene; ds.
   The present invention provides the protein and coding sequences o number of human reproductive system related antigens. These can be in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence enco protein of the invention
   21-JUN-2002
  ABL98314 standard; DNA; 31474
   Sequence
               WO200155317-A2
                                     Homo sapiens.
   ABL98314;
   Disclosure;
  Rosen
   (HUMA-)
   3073
  2001-465570/50
   testicular antigen
   Š
   72
   l Similarity
50; Conserv
  d nucleic acid molecule encoding a reproductive preventing, treating or ameliorating a medical
   HUMAN
  31474 BP; 9245
   AGATTGTGCCACTGCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   Barash SC,
  SEQ ID NO 8149; 1297pp + Sequence Listing; English.
  2000US-0249215P.
2000US-0249215P.
2000US-0249211P.
2000US-0249214P.
2000US-0249244P.
2000US-0249244P.
2000US-0249249P.
2000US-0249264P.
2000US-024929P.
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2000US-025039P.
2000US-0251930P.
2000US-0251988P.
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2000US-0251868P.
2000US-0251868P.
2000US-0251868P.
2000US-0251868P.
2000US-0251869P.
2000US-0251989P.
2000US-0251999P.
2000US-0251999P.
2000US-0251999P.
2000US-0251999P.
2000US-0251999P.
2000US-025499P.
  Conservative
   (first entry)
   GENOME
  1.6%; or-
100.0%; Pr
  SCI
  Ruben SM
   ₽,
  encoding
   6055
  Score 50;
Pred. No.
   ВP
   ü
  Mismatches
  DNA
   6292
  fragment
  DB 4; Le
1.3e-12;
   G;
   9882
  0
  SEQ
  Length 31474;
   T; 0
  Indels
  IJ
   U; O
  ö
  system ant condition.
   Other;
  3122
   23
  0;
  encoding
  antigen
  Gaps
   be of
   used
  a
  þ
   18
```

01-SEP-2000
01-SEP-2000
05-SEP-2000
06-SEP-2000
06-SEP-2000
08-SEP-2000
08-SEP-2000
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27-SEP-2000
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29-SEP-2000
20-CCT-2000
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08-NCV-2000
08-NCV-2000
08-NCV-2000
08-NCV-2000
08-NCV-2000
08-NCV-2000
08-NCV-2000
01-NCV-2000

2000US-022934AP.
2000US-0229513P.
2000US-0229513P.
2000US-023943P.
2000US-023124AP.
2000US-023124AP.
2000US-0231413P.
2000US-0231414P.
2000US-0231414P.
2000US-0232081P.
2000US-0232081P.
2000US-0232398P.
2000US-0232399P.
2000US-0232399P.
2000US-0232399P.
2000US-0232399P.
2000US-0233239P.
2000US-0233239P.
2000US-0233239P.
2000US-0233239P.
2000US-0233239P.
2000US-0233239P.
2000US-0234274P.
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2000US

| 17-JAN-2000; 2001WO-US001329. 131-JAN-2000; 2000US-0180628P. 04-FEB-2000; 2000US-0180628P. 14-FEB-2000; 2000US-0180628P. 16-MAR-2000; 2000US-0198123P. 11-MAR-2000; 2000US-0198123P. 11-MAR-2000; 2000US-0198123P. 11-MAR-2000; 2000US-0215133P. 11-JUN-2000; 2000US-0216880P. 11-JUN-2000; 2000US-0216880P. 11-JUN-2000; 2000US-0216880P. 11-JUN-2000; 2000US-0216847P. 11-JUN-2000; 2000US-021748PP. 11-JUN-2000; 2000US-0218290P. 14-JUG-2000; 2000US-022513P. 14-JUG-2000; 2000US-022575P. 14-JUG-2000; 2000US-0231414P. 16-JUG-2000; 2000US-023                                                                                                                                                                                                                                                                                                                         | 02-AUG-2001. |
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| 2000US-01390<br>2000US-01390<br>2000US-01390<br>2000US-01846<br>2000US-01846<br>2000US-01991<br>2000US-01991<br>2000US-02166<br>2000US-02166<br>2000US-02166<br>2000US-02166<br>2000US-02166<br>2000US-02166<br>2000US-02166<br>2000US-02166<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0226<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0231<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>2000US-0233<br>200 |              |
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| \\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | PR           |
| 29-SEP-2000; 22-SEP-2000; 22-CCT-2000; 22-CCT-2000; 22-CCT-2000; 22-CCT-2000; 23-CCT-2000; 23-CCT-2000; 20-CCT-2000; 20-NOV-2000; 21-NOV-2000; 21-NO                                                                                                                                                                                                                                                                                                                         | -SEP-2000;   |
| 07 2000US-0236379P. 07 2000US-0236379P. 07 2000US-0237037P. 07 2000US-0237039P. 07 2000US-0237039P. 07 2000US-023933P. 07 2000US-023933P. 07 2000US-0249990P. 07 2000US-0241785P. 07 2000US-0241786P. 07 2000US-0241809P. 07 2000US-0246477P. 07 2000US-0246478P. 07 2000US-0246532P. 07 2000US-0249211P. 07 2000U                                                                                                                                                                                                                                                                                                                         | 2000US-0     |

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RESULT 50
AAS30115/C
ID AAS30115/C
XX AAS30
XX AAS30
DT 21-NC
XX Lung
CH Chick
XW Chick
XW Cereb
XW Cereb
XW Santix
XW Santix
XW Wound
XW Hypex
XW Wound
XW Homo
XX Homo
XX Homo
XX 10-AU
PP 17-JP
XX 2-PE
PR 02-PE
PR 11-JP
PR 11-M
PR 11-M
PR 11-M
PR 11-J-M
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02-MAR-2000

16-MAR-2000

11-MAR-2000

11-MAR-2000

19-MAY-2000

07-JUN-2000

28-JUN-2000

30-JUN-2000

07-JUL-2000

07-JUL-2000

07-JUL-2000

11-JUL-2000

11-JUL-2000

11-JUL-2000
  Lung antigen protein; human; mouse; rabbit; goat; horse; cat; dog; chicken; sheep; immunosuppressive; antiarthritic; vasctropic; antirheumatic; antiproliferative; cytostatic; cardiant; neuroprotective; cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer; ophthalmological; vulnerary; gene therapy; autoimmune disease; neoplasm; hyperproliferative disorder; breast; liver; cardiovascular disorder; ds; cerebrovascular disorder; nervous system disorder; bacterial infection; fungal infection; viral infection; ocular disorder; endocrine disorder; gastrointestinal disorder; renal disorder; respiratory disorder; wound healing; skin aging; organ transplantation; food preservative; tissue regeneration; anti-infertility; food additive.
  0115/c
AAS30115
  The present invention provides the protein and coding sequences of 973 human testicular antigens, and fragments of their genomic sequences. The sequences can be used in the treatment of cardiovascular, urinary system, reproductive system, immune, respiratory, neurological and gastrointestinal disorders, infections, and particularly cancer, especially testicular cancers. The present sequence is a DNA encoding a
   02-AUG-2001
   WO200155303-A2
  Human
  21-NOV-2001
  AAS30115;
   Sequence 31474
   Disclosure; SEQ ID
   Nucleic acids encoding 973 human testicular antigen polypeptides, useful for preventing, diagnosing and/or treating testicular cancer.
  17-JAN-2001;
  3073
   sapiens.
   lung
  72
   50;
  Similarity
   fragment of the invention
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  standard;
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 23
   antigen genomic DNA #185
; 2000US-0179065P,
2000US-0184664P,
2000US-0184654P,
2000US-0186350P,
2000US-0198074P,
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2000US-0205515P,
2000US-0205467P,
2000US-021486F,
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   2001WO-US001301.
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   NO 2966; 766pp; English.
  DNA;
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  1.6%;
  32189
  6055 C;
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  뫄
   6292
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J. 1.3e-12;
O;
  Ģ
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    18-AUG 2000
22-AUG 2000
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23-AUG 2000
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01-SEP 2000
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2000US-0224519P.
2000US-0225213P.
2000US-0225214P.
2000US-0225266P.
2000US-0225267P.
2000US-0225268P.
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08-NOV-2000
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08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
08-NOV-2000
17-NOV-2000
17-NOV
              Sequences AAS29931-AAS30164 represent genomic DNA molecules, which encode the lung antigen polypeptides of the invention. Lung antigen polypeptides and their associated polypeptides are useful in the diagnosis, treatment and prevention of various types of disorders in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A pathological condition can be determined by detecting the presence or absence of a mutation in a lung antigen polymucleotide. The treatable disorders include autoimmune diseases such as rheumatoid arthritis, hyperproliferative disorders such as neoplasms of the breast or liver, cardiovascular disorders such as cardiac arrest, cerebrovascular disorders such as cardiac arrest, viruses such as alienter's disease, infections caused by bacteria, viruses such as premature labour and infertility, gastrointestinal disorders such as premature labour and infertility, gastrointestinal disorders such as premature labour and infertility, gastrointestinal disorders such as content of the premature of the premature is and the premature of the premature o
premature
Crohn's di
   Claim 1;
   Isolated
   WPI; 2001-457723/49.
  Rosen
   (HUMA-) HUMAN GENOME SCI
  testing and detection e.g. diagnosis.
  Ç
   polypeptide for treating, preventing and/ or prognosing ory disorders related to the lung including lung cancers
   SEQ ID NO 379; 507pp; English.
  Barash
  2000US-0246478

2000US-0246523P

2000US-0246525P

2000US-0246525P

2000US-0246525P

2000US-0246527P

2000US-0246528P

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2000US-0249245P.
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2000US-0249265P.
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2000US-0249269P.
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2000US-0251990P
2000US-0254097P
   2000US-0246477P
  SC,
  Ruben
  S
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renal disorders

such as glomerulonephritis

and also

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RESULT 51
RDB33452/c
ID ADB33452/c
ID ADB33452/c
ID ADB33452/c
AC ADB334
XX ADB33452/c
AC ADB334
XX ADB33452/c
AC ADB334
XX GAL
AC ADB34
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   Matches
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24-FBB-2000
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30-JUL-2000
07-JUL-2000
11-JUL-2000
11-JUL-2000
11-JUL-2000
14-JUL-2000
26-JUL-2000
14-JUL-2000
   respiratory disorders such as asthma and pleurisy. The polypeptides can also be used to aid wound healing, to prevent skin aging due to sunburn, to maintain organs before transplantation, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
   polyarteritis nodosa; bleeding disorder; thrombocytopenia; von Willebrand's disease; acquired platelet dysfunction; kidney failure; multiple myeloma; macrophage related disorder; Gaucher's disease; Neimann-Pick disease; tumour; colon cancer; pencreatic cancer; renal disorder; nephritis; bone disorder; Albers-Schonberg disease; bowleg; muscle disorder; Becker's muscular dystrophy; buchenne's muscular dystrophy; nervous disorder; ischaemic lesion; traumatic lesion; endocrine disorder; Cushing's syndrome; corticosteroid deficiency; gastrointestinal disorder; dysphagia; gastric reflux; human; ds.
  gene therapy; lung antigen; neoplasia; acute myelogenous leukaemia; adenocarcinoma; respiratory disorder; chronic rhinitis; sinusitis; immunodeficiency; X-linked agammaglobulinaemia; X-linked infantile agammaglobulinaemia; inflammatory disorder; adrenalitis; alveolitis; immune complex disease; serum sickness;
  Human novel lung related polypeptide DNA SEQ
  04-DEC-2003
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  20-MAR-2003
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llarity 100.0%;
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Pred. No.
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   Gaps
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1-AUG-2000; 1-AUG-2000; 1-AUG-2000; 1-AUG-2000; 1-AUG-2000; 1-AUG-2000;

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The invention relates to an isolated lung antigen polypeptide sequence or concoded sequence in a cDNA clone. The polypeptide and its polynucleotide are useful for treating, preventing, diagnosing and/or prognosing concepts and/or disorders such as pathological cell proliferative concepts as e.g. acute myelogenous leukaemias, adenocarcinoma; respiratory concepts such as chronic rhinitis, sinusitis; immunodeficiencies such as chronic rhinitis, sinusitis; immunodeficiencies such as concepts and concepts are concepts and co
  08-NOV-2000
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08-NOV-2000
17-NOV-2000
17-NOV
  Novel isolated lung antigen | diagnosing acute myelogenous Von Willebrand's disease.
   Disclosure;
  (HUMA-)
  Š
  2003-695900/66
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   Ruben
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  2000US.
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S-0246532P.
S-0246532P.
   NO 379; 178pp; English.
   SCI
   Barash
   polypeptides useful for treating, preventing, s leukemias, adenocarcinoma, thrombocytopenia,
   SC
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| FT exon FT intre FT exon FT intre FT exon FT exon FT exon FT exon FT exon FT intre FT exon FT exon FT exon FT intre FT exon FT exon FT intre FT exon FT exon FT exon FT intre FT exon FT intre FT exon | exon 4406. 4406. /*tag= intron /*tag= exon 6647. intron /*tag= intron /*tag= intron /*tag= exon /*tag= fitag= exon /*tag=                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | intron 2354. 2354. 2354. 2393. exon 2393. intron 2767. 2767. 2870. 414g= exon /*tag= intron /*tag= intron 3379. 414g= exon 339. 3462. 3950. |                     | in; gene roliferat diac arre sess; ner sess; ner on; ocula ood prese tive; wou heumatic; hromosome | RESULT 52 AAD16595/c ID AAD16595 standard; DNA; 32193 BP.  XX AC AAD16595; XX DT 19-NOV-2001 (first entry) XX DB Human novel protein-encoding gene 7, SEQ ID NO:37. | CC; endocrine disorders such as Cushing's syndrome, corticosteroid  Query Match  Query Match  Best Local Similarity 100.0%; Score 50; DB 10; Length 32189;  Best Local Similarity 100.0%; Pred. No. 1.3e-12;  Matches 50; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  Qy 3073 AGATTGTGCCACTCCAGCCTGGGCAACAGAGCAAGAGTCTGTCT |
|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------|---------------------|----------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | exon intressed i | exon intrintrintrintrintri                                                                                                                  | exon intr exon intr | exon intr exon intr intr exon intr intr                                                            | int exo                                                                                                                                                             |                                                                                                                                                                                                                                                                                                                                 |

exon

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Query Match
Best Local Similarity
Matches 50; Conserv
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04-FEB-2000

24-FEB-2000

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11-MAR-2000

11-MAR-2000

11-MAY-2000

11-MAY-2000

07-JUN-2000

28-JUN-2000

07-JUL-2000

07-JUL-2000

11-JUL-2000

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  02-AUG-2001.
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   exon
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   exon
   exon
   exon
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  intron
  intron
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  intron
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  intron
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  intron
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Conservative 0;
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2000US-0184664P.
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2000US-0225214P.
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bu
   .36224
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  bq
.36084
br
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bm
.32657
bn
.32844
bo
.34941
.35137
  .32275
bl
  .29991
bk
   .26539
bi
.29316
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bg
.26324
bh
  .26053
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   Gaps
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RESULT 53
AAL36258 E
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XX
AAL36258 E
XX
AC
AAL36258 E
XX
AC
DE Human musc
XX
Cytostatic
KW Cytostatic
KW antiallerg
KW vulnerary;
KW neurologic
KW musculosk
XX
OS Homo sapie
PN 02-AUG-200
XX
OZ-AUG-200
XX
OZ-AUG-200
XX
OZ-AUG-200
XX
OZ-AUG-200
XX
OZ-AUG-200
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PR 11-AUG-20
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04-FEB-2000
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11-AUG-2000
11-AUG
  17-JAN-2001;
  WO200155367-A1
  Homo sapiens.
  08-JAN-2002
  standard;
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밁
28855
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musculoskeletal system related polynucleotide SEQ
                        (first entry)
  DNA;
   32193
   쁌
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Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder; neurological disease; infection; human; secreted protein; musculoskeletal system; ds.

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2000US-0229343P.
2000US-023943P.
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2000US-023943P.
2000US-023943P. 2001WO-US001338

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2000US-0246611P.
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2000US-0246611P.
2000US-024652P.
2000US-0246611P.
20
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(ABB03087-ABB04109) associated with the musculoskeletal system useful for CC (ABB03087-ABB04109) associated with the musculoskeletal system useful for CC preventing, treating or ameliorating medical conditions e.g. by protein CC or gene therapy. The genes are isolated from a range of human tissues CC disclosed in the specification. The nucleic acids, proteins, antibodies CC and (ant) agonists are useful in the diagnosis, treatment and prevention CC of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the CC atrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, CC lung, or urogenital, (b) immune disorders e.g. Addison's disease, CC allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, CC diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid CC arthritis and ulcerative colltis; (c) cardiovascular disorders such as CC myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. CC cerebral anoxia and epilepsy; and (f) infectious diseases such as vital, control of the printed specification, but was CC obtained in electronic forms part of the printed specification, but was CC obtained in electronic forms to reservence.
  17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
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17-NOV-2000;
17-NOV-2000;
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06-DEC-2000;
08-DEC-2000;
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Sequence
  Example
   cancers and also for
   WPI; 2001-451937/48.
                                  ftp.wipo.int/pub/published_pct_sequences
  (HUMA-) HUMAN
   ξ
  2;
32193 BP;
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   Barash SC,
  7 2000US - 0249245P.
2000US - 0249264P.
2000US - 0249265P.
2000US - 0249297P.
2000US - 0249299P.
2000US - 0250319P.
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2000US - 0251868P.
2000US - 0251869P.
  GENOME
10182 A; 6701 C;
   for treating, preventing and/or prognosing the musculoskeletal system including muscultesting and detection e.g. diagnosis.
  SCI INC
   Ruben
   SM:
  6066
  G; 9244 T; 0 U; 0
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musculoskeletal

```
RESULT 54
ABX59246/c
ID ABX592
XX
AC ABX592
X7 26-FEB
XX
DT 26-FEB
XX
CDNA e
XX
KW Gene;
   片
   ફ
  Query Match
Best Local S
Matches 50
Gene; ss; musculoskeletal system antigen; cancer; metastasis;
                        cDNA encoding novel human musculoskeletal system antigen #1590.
   ABX59246 standard;
  26-FEB-2003
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; Pred. No. 1.3:
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14-AUG-
22-AUG-
  periodontal regeneration; tissue transport; bone graft; skin aging; keratinocyte growth; hair loss; melanocyte growth; cell proliferation; keratinocyte growth; organ transplant; cell differentiation; body height; weight; cell growth; organ transplant; cell differentiation; bissue; hair colour; eye colour; skin; percentage of adipose tissue; pigmentation; cosmetic surgery; metabolism; biorhythm; caricadic rhythm;
   14-AUG
   17-JAN-2001;
   10-OCT-2002
   US2002147140-A1
  Homo
  depression; tendency for violence; pain; reproductive capability; hormone level; endocrine level; appetite; libido; memory; stress; storage capability; fat content; lipid content; protein content; carbohydrate content; vitamin content; cofactor content;
  post-operative tissue repair; limb regeneration; neuronal growth; neurodegenerative disorder; Alzheimer's disease; Parkinson's disease; AIDS-related complex; chondrocyte growth; bone regeneration;
   cardiovascular condition; wound, post-operative tissue repair; 1:
  re-vascularisation; thrombosis; arteriosclerosis; mineral
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  2000US-0226868P

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  2000US-0224518P.
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  content;
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AC ADG629
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DT 11-MAR
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DE Genomi
XX
KW neurop
KW gastro

(first entry)

ADG62943 standard; DNA; 32193 BP

Genomic DNA 11-MAR-2004

neuroprotective; nootropic; respiratory; cardiovascular; gastrointestinal; antiparkinsonian; immunosuppressive; d

dermatological;

encoding human NOVX protein seq id

밁 δ

28855 3073

AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 28806 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122 Matches

Local

Similarity

Conservative

0,

Mismatches

Indels

0,

Gaps

0

Pred. No.

1.3e-12 0;

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cc and limb regeneration; stimulates neuronal growth; can treat and prevent concurring in certain disorders or neurodegenerative conditions, such as, Alzheimer's disease, Parkinson's disease, and AIDS-crelated complex; stimulates chondrocyte growth, thus they can be used to complex prevents skin aging due to sunburn by stimulating combane bone and periodontal regeneration and aid in tissue transports or combane bone and periodontal regeneration and aid in tissue transports or combane prevents skin aging due to sunburn by stimulating combane prevents hair loss, since FGF family members cativate hair-forming cells and promotes melanocyte growth; stimulates cativate hair-forming cells and promotes melanocyte growth; stimulates cativate hair-forming cells and promotes melanocyte growth; stimulates cativate hair combination with other cytokines; maintains organs before transplantation or for supporting cell culture of primary tissues; cinduces tissue of mesodermal origin to differentiation of embryonic stem used in combination with other cytokines; maintains organs before concreases or decreases the differentiation or proliferation of embryonic stem cells, besides, haematopoietic lineage; modulates mammalian characteristics, such as, body height, weight, hair colour, eye colour, catem cells, besides, body height, weight, hair colour, eye colour, cosmetic surgery); modulates mammalian metabolism; changes mammal's metal consension, tendency for violence, tolerance for pain, reproductive capabilities, hormonal or endocrine levels, appetite, libido, memory, or stress; increases or decreases storage capabilities, fat content, lipid, components. This sequence encodes a novel human musculoskeletal system contents used in cated appetite and of the printed specification, but was obtained in electronic format directly contents of the printed specification, but was obtained in electronic format directly contents.
                    Query Match
  printed specification, but was obtained in electronic from the US patent office at ftp.seqdata.uspto.gov/sequence.html?DocID=20020147140
  (ROSE/)
(RUBE/)
(BARA/)
  Sequence 32193 BP; 10182 A; 6701 C; 6066 G; 9244 T; 0 U; 0
  sequence encoding musculoskeletal system associated polypeptides useful for detecting disorders, e.g., cancer or cancer metastases, in animals humans. The nucleic acid: stimulates re-vascularisation of ischaemic
  Disclosure; SEQ ID NO 2623; 321pp; English.
   Isolated nucleic acid molecules associated polypeptides, useful
  Rosen CA, Ruben SM,
  and other cardiovascular conditions; treats wounds due
   tissues associated with conditions such as thrombosis, arteriosclerosis,
   WPI; 2003-128199/12
   08-DEC-2000; 2000US-0251869P.
   invention describes an isolated nucleic acid
  post-operative tissue repair, and ulcers; stimulates angiogenesis
   RUBEN S
  BARASH S C
  ₹ ≥
1.6%;
   Barash
                    Score 50;
   SC
   encoding musculoskeletal for detecting disorders,
                       DB 8;
                    Length 32193;
   molecule comprising
  to injuries,
   e.g. cancer.
   Other;
   유
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| PR 24 FRB 2000; 2000US -018668P; PR 24 FRB 2000; 2000US -018668P; PR 24 FRB 2000; 2000US -018974P; PR 16 FMR 2000; 2000US -018974P; PR 16 FMR 2000; 2000US -018974P; PR 17 FMR 2000; 2000US -018974P; PR 18 FMR 2000; 2000US -0205515P; PR 18 FMR 2000; 2000US -0205515P; PR 27 FMR 2000; 2000US -0205647P; PR 27 FMR 2000; 2000US -021868P; PR 26 FMR 2000; 2000US -0218999; PR 21 FMR 2000; 2000US -0218999; PR 20 FMR 2000; 2000US -0218999; PMR 20 FMR 2000; 2000US -02 | antiinflammatory; antirheum antidiabetic; hepatotropic; antiarteriosclerotic; cardifungicide; gynaecological; immune system disorder; mus gastrointestinal disorder; renal disorder; proliferati systemic lupus erythematosu thyroiditis; anaemia; Grave allergy; nephritis; parkins atherosclerosis; myocardial NOVX.  Homo sapiens.  US2003207285-A1.  06-NOV-2003.  12-AUG-2002; 2002US-0021646                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |
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|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | ק<br>קר קר ק                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |
| 02-OCT-2000) 13-OCT-2000) 13-OCT-2000) 13-OCT-2000) 20-OCT-2000) 20-OCT-2000) 20-OCT-2000) 20-OCT-2000) 20-OCT-2000) 20-OCT-2000) 20-OCT-2000) 08-NOV-2000) 08-NOV-2000) 08-NOV-2000) 08-NOV-2000) 08-NOV-2000) 08-NOV-2000) 08-NOV-2000) 08-NOV-2000) 08-NOV-2000) 108-NOV-2000) 108-NOV-2000) 108-NOV-2000) 117-NOV-2000) 117-NOV-2000]                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         |
| 2000US-0237038P. 2000US-0237038P. 2000US-0237039P. 2000US-0237040P. 2000US-0239937P. 2000US-0241786P. 2000US-0241809P. 2000US-0241809P. 2000US-0244617P. 2000US-0246475P. 2000US-0246475P. 2000US-0246478P. 2000US-0246678P. 2000US-0246678P. 2000US-0246678P. 2000US-0246525P. 2000US-0246525P. 2000US-0246525P. 2000US-0246525P. 2000US-0246525P. 2000US-0246529P. 2000US-0246529P. 2000US-024651P. 2000US-0246611P. 2000US-0246611P. 2000US-0246611P. 2000US-0246611P. 2000US-0246611P. 2000US-024921P. 2000US-024929P.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     | 0232399<br>0232398<br>0232490<br>0232490<br>0233401<br>02333063<br>02333063<br>023423<br>023423<br>023423<br>023423<br>023423<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>02343<br>0234<br>0234 |

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ADJ29996/c
ID ADJ299
XX
AC ADJ299
AC ADJ299
XX
DT 20-MAY
XX
DE Human
XX
KW muscul
KW gene t
XX
OS Homo s
XX
PN US2004
XX
XX
N US2004
XX
XX
N 15-JAN
  δ
   밁
   CC protein comprising a sequence that is at least 95% identical to: a

CC polynucleotide fragment of any of the nucleotide sequences listed in the

CC specification, or of the cDNA sequences listed in the specification,

CC which is hybridisable to the nucleotide sequences; a polynucleotide

CC encoding a polypeptide or a polypeptide fragment, domain or epitope of

CC any of the amino acid sequences listed in the specification, or a

CC polypeptide or a polypeptide fragment, domain or epitope of

CC convant of the nucleotide sequences listed in the specification, or a

CC polynucleotide which encodes a species homologue of the above amino acid

CC sequences; or a polynucleotide sequences listed in the specification; a

CC polynucleotide which encodes a species homologue of the above amino acid

CC sequences; or a polynucleotide capable of hybridising under stringent

CC conditions to any of the above polynucleotides, where the polynucleotide

CC does not hybridise under stringent conditions to a nucleic acid molecule

CC molecule and polypeptide are useful in diagnosing, preventing, prognosing

CC molecule and polypeptide are useful in diagnosing, preventing, prognosing

CC and/or activity of the above polypeptide, such as neural disorders,

immune system disorders, muscular disorders, reproductive disorders,
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Best Local S
Matches 50
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05-DEC-2000;

05-DEC-2000;

06-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

08-DEC-2000;

11-DEC-2000;

11-DEC-2000;

11-DEC-2000;

11-DEC-2000;

11-DEC-2000;
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  New
   US2004009488-A1
  musculoskeletal system; cytostatic; osteopathic; cancer; osteoporosis; gene therapy; vaccine; human; ds.
   Human musculoskeletal system-associated genomic DNA - SEQ ID 2623
   20-MAY-2004
  ADJ29996;
  ADJ29996 standard; DNA; 32193 BP
   Disclosure; SEQ ID NO 37; 194pp; English.
  treating disc
polypeptide,
   WPI; 2003-901052/82.
   Rosen
                       15-JAN-2004
  (HUMA-) HUMAN GENOME SCI INC.
   56
   polypeptides and nucleic acid molecules for diagnosing, preventing on atting diseases associated with aberrant expression or activity of the ypeptide, e.g. cancer, asthma, AIDS, Parkinson's disease or diabetes.
  3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   invention
  sapiens
   Ş
  50;
  Similarity
   Ruben SM,
  1.6%;
llarity 100.0%;
Conservative (
  ; 2000US-0251030P.

2000US-0251988P.

2000US-0256719P.

; 2000US-0251479P.

; 2000US-0251868P.

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; 2000US-0254997P.

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; 2001US-0254997P.
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   describes an isolated nucleic acid molecule (I) encoding a ising a sequence that is at least 95% identical to: a
   Barash
  0
  Score 50; pred. No.
   SS
  Mismatches
  DB 10;
  1.3e-12;
  <u>.</u>
  Length 32193;
  Indels
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  Gaps
  0
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118-AUG-2000
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   28-JUN-2000;
30-JUN-2000;
07-JUL-2000;
07-JUL-2000;
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14-AUG-2000;
  07-JUN-2000;
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2000US-018466P

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2000US-0198129P

2000US-0215135P

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2000US-0216880PP

2000US-0216880PP

2000US-0218290PP

2000US-0224518PP

2000US-02252613PP

2000US-02252613PP

2000US-02252613PP

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2000US-0233493PP

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06-DEC-2000
08-DEC-2000
  Rosen CA,
2004-090458/09
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  Ruben SM,
  2000US-0240960P

2000US-0241785P

2000US-0241788P

2000US-0241809P

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2000US-0246525P

2000US-0246525P

2000US-0246525P

2000US-0246528P

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2000US-0246510P

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2000US-0246510P

2000US-0249211P

  2000US-0251856P.
2000US-0251868P.
2000US-0251869P.
  2000US-0236370P
  2000US-0254097P
  GENOME
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31-JAN-2000 04-FEB-2000 24-FEB-2000 02-MAR-2000 16-MAR-2000 11-MAR-2000 18-APR-2000 19-MAY-2000 07-JUN-2000 28-JUN-2000 28-JUN-2000 07-JUL-2000 07-JUL-2000 11-JUL-2000 11-JUL-2000 11-JUL-2000 11-JUL-2000

2000US-0186350P. 2000US-0189874P.

2000US-0190076P.
2000US-0198123P.
2000US-0205515P.
2000US-0214886P.
2000US-0214886P.
2000US-0216847P.
2000US-0216880P.
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2000US-0216880P.

17-JAN-2001;

2001WO-US001301.

02-AUG-2001. WO200155303-A2 Homo sapiens.

A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER A SER

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RESULT 57
AAS30113/c
   FFFX8X55555555555XX
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  S
  Query Match
Best Local S
Matches 50
   The invention relates to a novel isolated musculoskeletal system-
associated nucleic acid molecule. The nucleic acid of the invention
demonstrates cytostatic and osteopathic activities and may be useful for
preparing a medicament for preventing, treating or ameliorating a medical
condition such as cancer of the musculoskeletal tissues or osteoporosis,
possibly via gene therapy or vaccine production. The current sequence is
that of the human musculoskeletal system-associated genomic DNA of the
invention. The current sequence is not shown within the specification per
se but is available on the USPTO web-site
   Lung antigen protein; human; mouse; rabbit; goat; horse; cat; dog; chicken; sheep; immunosuppressive; antiarthritic; vasotropic; antirheumatic; antiproliferative; cyrostatic; cardiant; neuroprotective; cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer; ophthalmological; vulnerary; gene therapy; autoimmune disease; neoplasm; hyperproliferative disorder; bracest; liver; cardiovascular disorder; decrebrovascular disorder; servous system disorder; bacterial infection; fungal infection; viral infection, ocular disorder; endocrine disorder; gastrointestinal disorder; renal disorder; respiratory disorder; gastrointestinal disorder; renal disorder; respiratory disorder; tissue regeneration; anti-infertility; food additive.
  AAS30113
  Sequence
   Disclosure; SEQ ID NO 2623; 289pp; English.
   preventing, treamusculoskeletal
  AAS30113;
   http:seqdata.uspto.gov/sequence.html?DocID=20040009488.
   Human lung antigen genomic DNA #183.
   21-NOV-2001
  28855
  nucleic acid molecule, useful for preparing a medicament for venting, treating or ameliorating a medical condition e.g., cancer
  3073
  l Similarity
50; Conserv
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   32193 BP;
   standard;
  Conservative
   (first entry)
   tissues or osteoporosis.
   DNA;
  10182 A; 6701 C; 6066 G; 9244 T; 0 U;
   1.6%;
  32221
  0;
   Score 50;
Pred. No.
   ВP
   Mismatches
  DB 12;
1.3e-12
  .3e-12;
  Length
   Indels
   0 Other;
  28806
  3122
  0
  Gaps
   0f
   0
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08-NOV-2000
017-NOV-2000
117-NOV-2000
Sequences AAS29931-AAS30164 represent genomic DNA molecules, which enc the lung antigen polypeptides of the invention. Lung antigen polypeptil and their associated polynucleotides are useful in the diagnosis, treatment and prevention of various types of disorders in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A pathological condition can be determined by detecting the presence or absence of a mutation in a lung antigen polynucleotide. The treatable disorders include autoimmune diseases such as rheumatoid arthritis, hyperproliferative disorders such as neoplasms of the breast or liver, cardiovascular disorders such as cardiac arrest, cerebrovascular disorders such as cardiac arrest, cerebrovascular disorders such as cerebral ischaemia, nervous system disorders such as Alzheimer's disease, infections caused by bacteria, viruses and fungi,
  Isolated polypeptide for treating, preventing and/or prognosing respiratory disorders related to the lung including lung cancers for testing and detection e.g. diagnosis.
   Rosen
  Claim
  (HUMA-)
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   HUMAN
   SEQ
  Barash
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20
   GENOME
  ð
  SC,
   377;
   3-0246475P

3-0246477P

3-0246477P

3-0246523P

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3-0246523P

3-0246523P

3-0246532P

3-0246532P

3-0246532P

3-0249211P

3-0249219P

3-025919P

3-025919P

3-025919P

3-0251186P

3-0251186
   SCI
  Ruben
  507pp; English.
   MS
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which encode polypeptides

and also

14-JUL-2000
26-JUL-2000
14-AUG-2000
12-AUG-2000
12-AUG-2000
13-AUG-2000
13-AUG-2000
13-AUG-2000
10-SEP-2000
01-SEP-2000
02-OCT-2000

2000US-0218290P 2000US-022964P 2000US-0225266P 2000US-0225266P 2000US-0225270P 2000US-0225270P 2000US-0225270P 2000US-0225758P 2000US-0231243P 2000US-0241785P 2000US-0241808P 2000US-0241808P 2000US-0241808P 2000US-0241826P

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RESULT 58
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ID
ADB33450 standard; DNA; 32221 BP
XX
AC
ADB33450;
XX
C
ADB33450;
XX
DF
O4-DEC-2003 (first entry)
XX
DE
Human novel lung related polypep
XX
DE
Human novel lung antigen; neop
XX
DE
Human novel lung antigen; neop
XX
DE
KW
Gene therapy; lung antigen; neop
XX
VInked infantile agammaglobuli
XX
VInked infantile agammaglobuli
XX
VON Willebrand's disease; acquir
XX
VON Wille
  88888888888
  밁
  8
  Matches
   Query Match
Best Local Similarity
   ocular disorders such as corneal infection, endocrine disorders such as premature labour and infertility, gastrointestinal disorders such as Crohn's disease, renal disorders such as glomerulonephritis and respiratory disorders such as asthma and pleurisy. The polypeptides can also be used to aid wound healing, to prevent skin aging due to sunburn, to maintain organs before transplantation, to regenerate tissues and in chemotaxis. The polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
  immunodeficiency; X linked agammaglobulinaemia;
X-linked infantile agammaglobulinaemia; inflammatory disorder;
adrenalitis; alveolitis; immune complex disease; serum sickness;
polyarteritis nodosa; bleeding disorder; thrombocytopenia;
Von Willebrand's disease; acquired platelet dysfunction; kidney fai:
multiple myeloma; macrophage related disorder; daucher's disease;
weimann-Pick disease; tumour; colon cancer; pancreatic cancer;
renal disorder; nephritis; bone disorder; Albers-Schonberg disease;
bowleg; muscle disorder; Becker's muscular dystrophy;
Duchenne's muscular dystrophy; nervous disorder; ischaemic lesion;
traumatic lesion; endocrine disorder; Cushing's syndrome;
traumatic lesion; endocrine disorder; Cushing's syndrome;
   gene therapy; lung antigen; neoplasia; acute myelogenous leukaemia;
   Human novel lung related polypeptide DNA SEQ ID NO 377.
  12548
  50;
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 12499
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  1.6%;
ilarity 100.0%;
Conservative (
   deficiency; gastrointestinal
  k; Score 50; DB
k; Pred. No. 1.3
0; Mismatches
  disorder;
   ₽₽
  chronic
  DB 5;
   1.3e-
   disorder; dysphagia;
  12;
  rhinitis; sinusitis;
  Length 32221;
  Indels
   kidney failure;
  0;
  Gaps
  0
       14-AUG-2000
14-AUG-2000
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22-AUG-2000
22-AUG-2000
23-AUG-2000
01-SEP-2000
02-OCT-2000
03-NOV-2000
08-NOV-2000
08-NOV-2000
   2000US-0237040P.
2000US-0239935P.
2000US-0239937P.
2000US-0240960P.
2000US-0241221P.
  2000US-0226868P.
2000US-0227182P.
2000US-0227009P.
2000US-0228924P.
2000US-0228924P.
2000US-0229344P.
   2000US-0224518P.
2000US-0225214P.
2000US-0225214P.
2000US-0225266P.
2000US-0225268P.
2000US-0225270P.
2000US-022547P.
2000US-022547P.
2000US-0225479P.
2000US-02257579.
2000US-022575P.
2000US-022579P.
2000US-022579P.
2000US-022579P.
   2000US-0241785P.
2000US-0241786P.
2000US-0241780P.
2000US-0241809P.
2000US-0241829P.
2000US-0244617P.
2000US-0246475P.
2000US-0246475P.
2000US-0246476P.
   2000US-0233064P.
2000US-0233065P.
2000US-0234221P.
2000US-0234274P.
2000US-023499PP.
2000US-023499PP.
2000US-023494P.
  2000US-0229345P
2000US-0229519P
2000US-0230437P
2000US-0231243P
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2000US-02312443P
2000US-02312443P
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2000US-0232981P
2000US-0232981P
2000US-0232399P
  2000US-0237038P.
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   2000US-0237
2000US-0237
   2000US-02363
  2000US-0235
   2000US-02368
  2000US-02363
   200008-02363
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08-NOV-2000

17-NOV-2000

17-NOV-2000

17-NOV-2000
  17-NOV-2000;
11-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
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08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
  17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
The invention relates to an isolated lung antigen polypeptide sequence or encoded sequence in a cDNA clone. The polypeptide and its polypucleotide are useful for treating, preventing, diagnosing and/or prognosing diseases and/or disorders such as pathological cell proliferative neoplasias e.g. acute myelogenous leukaemias, adenocarcinoma; respiratory disorders such as chronic rhinitis, sinusitis; immunodeficiencies such as x-linked agammaglobulinaemia; x-linked agammaglobulinaemia; inflammatory disorders such as adrenalitis, alveolitis; immune complex diseases such as serum sickness, polyarteritis nodosa; bleeding disorders such as thrombocytopenia, Von Willebrand's disease; acquired platelet dysfunction such as kidney failure, multiple myeloma; disorders associated with macrophage numbers and/or macrophage function such as schmann-pick disease; tumours such as colon cancer, pancreatic cancer; renal disorders such as kidney failure, nephritis;
  17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
  WPI;
  Disclosure; SEQ ID NO 377; 178pp; English
  Novel isolated lung antigen polypeptides useful for treating, preventing, diagnosing acute myelogenous leukemias, adenocarcinoma, thrombocytopenia, Von Willebrand's disease.
   17-NOV-2000;
17-NOV-2000;
   17-NOV-2000;
  2003-695900/66
   Ş
  HUMAN
   Ruben SM,
   2000US-0246528P.
2000US-0246632P.
2000US-0246610P.
2000US-0246611P.
2000US-0249208P.
2000US-0249209P.
2000US-0249210P.
2000US-0249211P.
2000US-0249211P.
2000US-0249213P.
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2000US-0249213P.
2000US-0249213P.
2000US-0249214P.
  GENOME
  SCI INC
   Barash
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닭
  Ś
   8888
   Matches 50;
  Query Match
  Best Local
  The invention relates to a method for identifying a molecule involved in lipid regulation comprising identifying a molecule that binds to or inhibits binding of a molecule to high bone mass (HBM) or its wild type gene, Zmax1. Compounds identified by the method are useful for treating, diagnosing, preventing or screening for normal and abnormal lipid-associated conditions, including arteriosclerosis, cardiovascular disease, stroke, and osteoporosis. The compounds may also be used in the treatment or prevention of diabetic atherosclerosis, neurovascular conditions caused by plaque build-up, poor clirculation due to plaque build-up and associated poor wound-healing. The methods may be used in gene therapy, pharmaceutical development, and diagnostic assays for bone development disorders. Molecules identified by comparison of Zmax1 and
   bone disorders such as Albers-Schonberg disease, bowlegs; muscle disorders such as Becker's muscular dystrophy, Duchenne's muscular dystrophy, nervous disorders such as ischaemic lesions, traumatic lesions; endocrine disorders such as Cushing's syndrome, corticosteroid
   Identifying molecules involved in lipid regulation, useful for diagnosing, treating or preventing e.g., arteriosclerosis, comgidentifying a molecule that binds to high bone mass gene or its corresponding wild type gene.
   lipid-associated condition; arteriosclerosis; cardiovascular disease; ss; osteoporosis; atherosclerosis; diabetic atherosclerosis; plaque build-up; neurovascular condition; wound healing; gene therapy; PCR primer; probe;
                     molecules encoding human Zmax1 and HBM, and PCR primers, and adapters of the invention
  HBM systems can be used as surrogate markers in pharmaceutical development, in diagnosis of human or animal bone disease, and in the treatment of bone diseases. Sequences ABK22776-ABK23411 represent cDN:
   Carulli JP,
   09-APR-2002
  Example 2; Page
  26-MAY-2000; 2000US-00578900
   25-MAY-2001; 2001WO-US016946.
  06-DEC-2001.
  osteopathic;
  bone development disorder; antiarteriosclerotic; cardiovascular;
   Human; mouse;
  Human high bone mass (HBM)
   ABK22783
  ABK22783
   12548
  3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   h 1.6%;
Similarity 100.0%;
  UNIV CREIGHTON SCHOOL MEDICINE
   GENOME THERAPEUTICS CORP.
  standard; cDNA; 36305
   Conservative
   Little RD,
  cerebroprotective.
   (first entry)
   Zmax1; HBM; high bone mass gene;
  323-350; 409pp;
   Recker RR, Johnson
  polymucleotide clone
   0
  Score 50;
Pred. No.
   Mismatches
  English
  1.3e-12;
  DB 10;
   lipid regulation; stroke;
  Length 32221;
   Indels
                                    probes, linkers
  or its
  comprises
   0
   Gaps
   0
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RESULT 60
ACN44230/c
RESULT 61
ACN44786
ID ACN44
   맑
   S
   S
   밁
  S
   The present invention relates to novel DNA and protein sequences which care associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of a bioactive agent capable of modulating the activity of CAP; (iv) for carcinoma; (vi) for inhibiting the activity of CAP; (vi) for treating carcinoma; (vii) for neutralizing the effect of CAP; (vii) for treating carcinoma; (viii) for neutralizing the effect of CAP; (ix) as a biochip; (cx) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining carcinoma or a propensity to carcinoma; (viii) for carcinoma associated (CA) gene copy number. In addition, the CAP genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent
   Query Match
Best Local 9
   Matches
  Matches
   Query Match
Best Local
   Sequence 36305 BP; 7938 A; 9658 C; 10106 G; 8602 T; 0 U; 1 Other;
  Human genomic sequence hCG21559.
   ACN44230 standard;
ACN44786 standard; DNA; 156843 BP.
   Recombinant nucleic acid useful for diagnosis and treatment of carcinoma
   Homo sapiens.
  Cytostatic; carcinoma; lymphoma; cancer; human; gene; ss.
  18-NOV-2004
  ACN44230;
   Sequence 66973
  Claim 1; SEQ ID NO 574; Opp; English.
  comprises a nucleotide sequence.
  WPI; 2003-328604/31.
   Morris DW
  01-MAR-2002; 2002US-00087192
  28-FEB-2003; 2003WO-US006235.
  12-SEP-2003
   WO2003073826-A2
  (SAGR-) SAGRES DISCOVERY.
  તે
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   5112 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 5161
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   Similarity
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100.0%; Pred. No.
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100.0%; Pr
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RESULT 62
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XX ADP75
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KW ADAMT
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XW ASTHMINAN
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XX HUMAN
XX HUM
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Best Local Similarity
   The present invention relates to novel DNA and protein sequences are useful for: (i) for are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of a bioactive agent capable of modulating the activity of CAP; (iv) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing carcinoma; (vi) for inhibiting the activity of CAP; (vii) for treating carcinoma; (viii) for neutralizing the effect of CAP; (vii) as a biochip; (x) for diagnosing carcinoma are appeared (CA) gene copy number. In addition, the determining Carcinoma appeared (CA) gene copy number. In addition, the screen of the capable of t
   determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent US2002182586A1, for which no sequence data was published
   Human; chromosome 5; ds; gene; ADAM19; Endophilin 1; Endophilin 2; NRG2; ADAM782; a disintegrin and metalloprotease; neuroregulin 2; SNP; single nucleotide polymorphism; a disintegrin and metalloprotease with thrombospondin typel motif 2; asthma; atopy; obesity; inflammatory bowel disease; respiratory disorder.
   Recombinant nucleic acid useful for diagnosis and treatment comprises a nucleotide sequence.
  01-MAR-2002; 2002US-00087192
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   Claim 1; SEQ ID NO 1408; Opp; English.
  WPI; 2003-328604/31.
   28-FEB-2003; 2003WO-US006235
   12-SEP-2003.
   WO2003073826-A2
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  Human ADAMTS2 gene.
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      Homo sapiens.
  12-AUG-2004
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  of carcinoma
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| 0 23                                                                                              | variation | 2 T T                                                   |                                                                                                  | 1.<br>1.<br>1.                          |
| single nucleoride                                                                                 | variation | F 17 1                                                  | .н ~~                                                                                            | variation                               |
| <pre>/standard name= "Single nucleotide polymorphism" replace(223251,T) /*tag= at</pre>           | variation | FT                                                      | . <b>H ~</b> ~ !                                                                                 | a i                                     |
| 6 5                                                                                               | variation | ગુગામ<br>મુખ્ય                                          | H ~~ F                                                                                           | variation                               |
| <pre>/*cay- au /standard_name= "Single nucleotide polymorphism" replace(221189,G) /*tag= ar</pre> | variation | 1 70 70 70 70 70 70 70 70 70 70 70 70 70                | /*tag= s<br>/*tag= s<br>/standard_name= "Single nucleotide polymorphism"                         | variation                               |
| e 2                                                                                               | variation | # TP                | · =                                                                                              | 1. 1                                    |
| @ B                                                                                               | variation | 77<br>77<br>77                                          | ı <b>~</b> ~ !                                                                                   | *************************************** |
| @ <b>B</b>                                                                                        | variation | 0 0 0 0 1                                               | 1 <b>~~</b> !                                                                                    | variation                               |
| @ B                                                                                               | variation | FT                                                      | . ~ ~                                                                                            | •                                       |
| <pre>/standard_name= "Single nucleotide polymorphism" replace(215294,A) /*tag= am</pre>           | variation | 27 T                                                    | н                                                                                                | variation                               |
| name= "Single nucleotide<br>[5293,T)                                                              | variation | 87<br>87<br>87                                          | .н ~~                                                                                            | variation                               |
| name= "Single nucleotide<br>[5171,A)                                                              | variation | eri<br>eri                                              | .н                                                                                               | variation                               |
| name= "Single nucleotide<br>[3555,A)<br>j                                                         | variation | । या या या<br>विस्तुति                                  | re *                                                                                             | variation                               |
| name= "Single nucleotide<br>13324,A)                                                              | variation | 1 T T T                                                 | H \ \                                                                                            | variation                               |
| name= "Single nucleotide<br>13294,A)                                                              | variation | 1 T T T T                                               | ч~~                                                                                              | variation                               |
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| /*E35- us<br>/standard_name= "Single nucleotide polymorphism"<br>replace(211462,T)<br>/*tag= af   | variation | 전 전 전 전<br>전 <b>전</b> 전 전 전 전 전 전 전 전 전 전 전 전 전 전 전 전 전 | /*tag= g<br>/*tandard_name=<br>replace(143623,                                                   | variation                               |
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| 0 22                                                                                              | variation | T T T T                                                 | replace (7805, C) /*tag= d /*tandard name=                                                       | : p                                     |
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   2000US-023935P;
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   GENOME SCI INC.
   Ruben SM;
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   S
  cc amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cc activity, and can be used in gene therapy and vaccine production. (I) cc proteins and polynucleotides may be used in the prevention, diagnosis and creatment of diseases associated with inappropriate (I) expression. For ce example, they may be used to treat disorders associated with decreased ce expression by rectifying mutations or deletions in a patient's genome ct that affect the activity of (I) by expressing inactive proteins or to complement the patients own production of (I). Additionally, (I) collected and be used to produce the secreted (I), by inserting the converse and treat immune/haematopoietic-related diseases, especially conneces and cancer metastases of haematopoietic articles peromic concers and cancer metastases of haematopoietic active peromic concers and cancer metastases of haematopoietic articles and AAM82169 concers from the present invention. AAK54942 to AAK54950 and AAM82169 corpresent sequences used in the exemplification of the present invention.
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CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic contivity, and can be used in gene therapy and vaccine production. (I) crotesins and polynucleotides may be used in the prevention, diagnosis and creatment of diseases associated with inappropriate (I) expression. For ceample, they may be used to treat disporders associated with decreased cexpression by rectifying mutations or deletions in a patient's genome cc that affect the activity of (I) by expressing inactive proteins or to concert affect the patients own production of (I). Additionally, (I) colored concert and bost cell and culturing the cell to express the cultic acids into a host cell and culturing the cell to express the complete and treat immune/haematopoietic-derived cells. AAK64703 cc ancers and cancer metastases of haematopoietic-derived cells. AAK64703 cc ancers and cancer metastases of haematopoietic aritigen genomic concerts are represent human immune/haematopoietic antigen genomic concerts and cancer metastases of haematopoietic antigen genomic concerts and haemat
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-----------|----------------------------------------------------------|-----------------------------|-----------------------------|----------|-------------|------------------|------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------|-----------------------------|-----------|------------------------------------------|-----------|-------------------------------------------------------|---------------------------------------------------------|--------|
| $egin{array}{c} 0.00000000000000000000000000000000000$                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 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                                                                                              | 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7                                                                                                                                                                                                                                         |                                                                              | בר<br>קק                                   | PR<br>PR                                                          | PR<br>PR                                   | PR<br>PR                                   | PR<br>PR                                   | PR                                         | אַק                                        | PR                                         | PR                  | PR                  | אק<br>אפ            | PR                  | PR                          | PR                                                         | PR                                                       | PR                          | קק<br>קק                    | מק<br>אק | PR          | PR<br>PR         | PR<br>PR                                                                                                                     | PR                                                                           | PR                          | PR<br>PR  | PR                                       | אק<br>אק  | PR<br>Rq<br>ac                                        | PR                                                      | מק     |
| $egin{array}{c} 0.00000000000000000000000000000000000$                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | 11111111111111111111111111111111111111                                                                                                                                                                                                                                        | 17-NOV<br>17-NOV<br>17-NOV                                                   | 08-NOV                                     | NON-80<br>NON-80                                                  | AON-80                                     | 08-NOV                                     | NON-80                                     | AON-80                                     | AON-80                                     | 08-NOV                                     | 20-001              | 20-0CI              | 20-0CI              | 20-0CI              | 13-001                      | 02-0CI                                                     | 02-001                                                   | 02-0CT                      | 29-SEP                      | 29-SEP   | 27-SEP      | 26-SEP<br>27-SEP | 25-SEP<br>25-SEP                                                                                                             | 21-SEP                                                                       | 14-SEP                      | 14-SEP    | 14-SEP                                   | 14-SEP    | 13 -80<br>138-80                                      | 08-SEP                                                  | 08-SEP |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 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                                                                                              | NON                                                                                                                                                                                                                                       |                                                                              |                                            |                                                                   |                                            |                                            |                                            |                                            |                                            |                                            |                     |                     |                     |                     |                             |                                                            |                                                          |                             |                             |          |             |                  |                                                                                                                              |                                                                              |                             |           |                                          |           |                                                       |                                                         |        |

```
RESULT 66
AAX86737C
ID AAX8677
XX AAX8677
XX AAX867
XX O7-NOV
DT 07-NOV
XX Y
XX Human
XX Human;
XW Human;
XW Human;
XX Cytost
XX OS Homo 8
   밁
  ঠ
  cc amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cc activity, and can be used in gene therapy and vaccine production. (I) croteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For cc example, they may be used to treat disorders associated with decreased cc expression by rectifying mutations or deletions in a patient's genome ct that affect the activity of (I) by expressing inactive proteins or to cc supplement the patients own production of (I). Additionally, (I) cc polynucleotides may be used to produce the secreted (I), by inserting the cc protein. (I) proteins and polynucleotides may be used to prevent, (I) cd isgnose and treat immune/haematopoietic-related diseases, especially cc cancers and cancer metastases of haematopoietic derived cells. AAK64703 cc antigen genomic sequences from the present invention. AAK54942 to AAK54950 and AAM82169 cc represent sequences used in the exemplification of the present invention.
   Query Match
Best Local S
Matches 49
   17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
17-NOV-2000
01-DEC-2000
01-DEC-2000
05-DEC-2000
05-DEC-2000
06-DEC-2000
06-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
                             Human; immune; haematopoietic; immune/haematopoietic antigen; cancer, cytostatic; gene therapy; vaccine; metastasis; ds.
   Sequence 272 BP;
  Nucleic
  Rosen
   Homo sapiens.
   Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41549.
   07-NOV-2001
   AAK86737;
   AAK86737 standard; DNA; 281 BP
   Disclosure; SEQ ID NO 41548; 3071pp + Sequence Listing; English
   (HUMA-) HUMAN GENOME
   Local Similarity
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT
  Ş
  64
  351 to AAK64702 encode the human immune/haematopoietic antigen acid sequences given in AAM82170 to AAM91921. (I) have cytosta
   49;
   c acids encoding human immune/hematopoietic antigen polypeptides, for preventing, diagnosing and/or treating cancers and metastasis.
  AGATTGTGCCACTGCAGCCCTGGGCAACAGAGCAAGACTCTGTCT
  Barash SC,
  2000US-0249264P.
2000US-0249265P.
2000US-0249299P.
2000US-0250160P.
2000US-0250160P.
2000US-0251030P.
2000US-0251989P.
2000US-0251868P.
2000US-0251868P.
2000US-0251869P.
   1.6%; Score 49; ilarity 100.0%; Pred. No. Conservative 0; Mismatci
  (first entry)
   <u>4</u>5
   A; 77 C; 67 G; 83 T; 0 U;
   SCI INC
  Z.
   Mismatches
   DB 4;
3. 4.5e-12;
0;
   0 Other;
  Length 272;
   Indels
  16
   3121
   0,
   Gaps
   Ξ
   0
   17-MAR-2000
18-APR-2000
19-MAY-2000
29-JUN-2000
20-JUN-2000
30-JUN-2000
11-JUL-2000
11-JUL-2000
11-JUL-2000
14-AUG-2000
1-SEP-2000
1-SEP-2000
0-SEP-2000
  WO200157182-A2
  17-JAN-2001;
  2000US-0189874P

2000US-0199874P

2000US-0199076P

2000US-0199076P

2000US-0209467P

2000US-021648P

2000US-021648P

2000US-021648P

2000US-0217487P

2000US-0217487P

2000US-0217490P

2000US-021749P

2000US-0224519P

2000US-0225214P

2000US-0225214P

2000US-0225266P

2000US-0225268P

2000US-0225447P

2000US-0225447P

2000US-0225757P

2000US-0225757P

2000US-0225759P
               2000US-0232399P.
2000US-0232400P.
2000US-0233401P.
2000US-0233063P.
2000US-0233065P.
2000US-0233065P.
2000US-0234223P.
2000US-0234274P.
2000US-023498P.
2000US-023498P.
  2000US-0229287P.
2000US-0229343P.
2000US-0229345P.
2000US-0229519P.
2000US-0239519P.
2000US-0231243P.
2000US-0231244P.
2000US-0231244P.
2000US-0231413P.
2000US-0231413P.
2000US-0231413P.
2000US-0231413P.
2000US-0231413P.
2000US-0231413P.
2000US-0231413P.
2000US-023198P.
2000US-023198P.
   2000US-0225759P.
2000US-0226279P.
2000US-0226681P.
   2001WO-US001354
  2000US-0227009P.
2000US-0228924P.
   2000US-0226868P.
2000US-0227182P.
  2000US-0179065P
```

polypeptides, and metastasis.

 $\Xi$ 

```
17.NOV-2000
17.NOV-2000
17.NOV-2000
17.NOV-2000
17.NOV-2000
01.DEC-2000
01.DEC-2000
05.DEC-2000
06.DEC-2000
06.DEC-2000
08.DEC-2000
08.DEC-2000
08.DEC-2000
   27-SEP-2000
29-SEP-2000
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29-SEP-2000
29-SEP-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
02-OCT-2000
03-OCT-2000
03-NOV-2000
04-NOV-2000
08-NOV-2000
08-NOV
  17-NOV-2000;
17-NOV-2000;
                        HUMAN
  2000US-0241785p
2000US-0241785p
2000US-0241808p
2000US-0241808p
2000US-02446475p
2000US-0246477p
2000US-0246477p
2000US-0246477p
2000US-0246477p
2000US-0246525p
2000US-0246525p
2000US-0246525p
2000US-0246525p
2000US-0246525p
2000US-0246525p
2000US-0246525p
2000US-0246527p
2000US-0246527p
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2000US-0246527p
2000US-0249211p
2000US-025921865p
2000US-02511868p
2000US-02511868p
2000US-02511868p
2000US-02511868p
2000US-02511868p
2000US-02511868p
2000US-02511869p
2000US-02511869p
2000US-02511869p
2000US-0251989p
   2000US-0235836P.
2000US-0236367P.
2000US-0236367P.
2000US-0236368P.
2000US-0236369P.
2000US-0236376P.
2000US-0236370P.
2000US-0236802P.
   2000US-0237037P
2000US-0237038P
2000US-0237039P
2000US-0237040P
2000US-0239935P
2000US-0239937P
2000US-0239937P
                        GENOME
                        SCI
  RESULT 67
AAK66626
  밁
  S
   Query Match
Best Local Similarity
Matches 49; Conser
31-JAN-2000;
04-FEB-2000;
24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
18-AR-2000;
19-MAY-2000;
07-JUN-2000;
28-JUN-2000;
30-JUN-2000;
07-JUL-2000;
07-JUL-2000;
11-JUL-2000;
11-JUL-2000;
   amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopolistic-related diseases, especially
  cancers and cancer metastases of haematopoietic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoletic antigen genomic sequences from the present invention. AAK54942 to AAK5950 and AAM82169 represent sequences used in the exemplification of the present invention
   09-AUG-2001.
  WO200157182-A2.
   cytostatic;
   Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
  Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21438.
  06-NOV-2001 (first entry)
   AAK66626;
  AAK66626 standard; DNA; 21477
   Sequence 281 BP; 48 A; 79 C; 70 G; 84 T; 0 U; 0 Other;
  AAK54951 to AAK64702 encode the human immune/haematopoietic antigen
   Disclosure; SEQ ID NO 41549; 3071pp + Sequence Listing; English
   useful for
  Rosen
   17-JAN-2001;
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT
   sapiens.
  2001-483426/52.
  ξ,
   63 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT
   acids encoding for preventing,
  Barash
2000US-0179065P.
2000US-0184664P.
2000US-0184664P.
2000US-0186350P.
2000US-0198076P.
2000US-0199076P.
2000US-0205515P.
2000US-0205467P.
2000US-0215135P.
2000US-0215135P.
2000US-0215135P.
2000US-0215135P.
2000US-0215135P.
2000US-0215135P.
2000US-0215135P.
2000US-0215135P.
2000US-0215135P.
   gene therapy; vaccine; metastasis; ds.
  Conservative
   2001WO-US001354
  SC,
  1.6%;
   human immune/hematopoietic antigen diagnosing and/or treating cancers
  Ruben
  0;
  Score 49;
Pred. No.
  SM.
   BP.
  Mismatches
```

DB 4; Le . 4.5e-12; 0 Length Indels 281;

3121 15

ç,

Gaps

0

(HUMA-)

```
01.NOV-2000
08.NOV-2000
09.NOV-2000
09.NOV-2000
01.NOV-2000
01.NOV
AAKS4951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis an treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent,
  Nucleic
useful f
   Rosen
   Disclosure;
  (HUMA-)
  2001-483426/52
   Ş
  c acids encoding for preventing,
  HUMAN
   Barash
   QES
   2000US-024647FP

2000US-024647FP

2000US-024647FP

2000US-024647FP

2000US-0246525P

2000US-0246525P

2000US-0246526P

2000US-0246526P

2000US-0246526P

2000US-024651P

2000US-0246611P

2000US-024921P

2000US-025921P

2000US-025921P

2000US-025921P

2000US-025921P

2000US-025921P

2000US-025921P

2000US-025921P

2000US-025929P

2000US-025186P

   GENOME
   ij
   sc,
   ŏ
  SCI
   21438; 3071pp +
  human immune/hematopoietic diagnosing and/or treating
   Ruben
   NS.
   Sequence Listing; English
  antigen polypeptides, cancers and metastasis.
```

the

and

11-JUL-2000
14-JUL-2000
14-JUL-2000
14-JUL-2000
14-JUG-2000
12-JUG-2000
12-JUG-2000
13-JUG-2000
13-JUG-2000
14-SEP-2000
10-SEP-2000
01-SEP-2000
02-OCT-2000

2000US-0217496P.
2000US-021964P.
2000US-022961P.
2000US-0225261P.
2000US-0225261P.
2000US-0225261P.
2000US-0225261P.
2000US-0225261P.
2000US-0225261P.
2000US-0225261P.
2000US-0225759P.
2000US-0231414P.
2000US-0231414P.
2000US-0231243P.
2000US-0

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RESULT 68
ADZ13418/c
  associated (CA) nucleic acid, comprising two or more nucleic acid probes. CC The invention also relates to a peptide array comprising two or more compound isolated polypeptides encoded by a CA nucleic acid sequence, a compound cc that binds to a polypeptide, which is prepared by immunizing a host animal composition comprising the polypeptide or its fragment which cc binds to a polypeptide, which is prepared by immunizing a host animal composition comprising the polypeptide or its antigen binding fragment, a composition comprising the cc fragment and collecting cells from the host expressing antibodies against the antipody and a carrier, a method of screening for anticancer activity, a method of detecting a CA nucleic acid, a method of diagnosing cancer, a cc method of treating cancer and a method of inhibiting expression of a CA nucleic acid in a cell. The CA nucleic acids are useful for detecting CA cc mucleic acids. The antibody is useful for detecting the presence or cabeence of cancer cells in an individual which involves contacting cells from the individual with the antibody and detecting a complex of a CA concert cells in an individual which involves contacting cells of the complex of a capture to the cancer cells and the antibody, where the detection of the complex correlates with the researce of cancer cells in an individual which involves contacting cells in the cancer cells and the antibody, where the detection of
   Matches 49;
  Query Match
Best Local
   Nucleic acid array useful for detecting cancer associated nucleic acid, comprises two or more nucleic acid probes.
  Morris DW, Malandro MS
  Homo sapiens
   Human cancer-associated genomic DNA #80.
  ADZ13418 standard;
   Sequence 21477 BP; 5311 A; 4999 C; 5256 G; 5911 T; 0 U; 0 Other;
   diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAKS4942 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention
the complex correlates with the presence of cancer cells in the individual. The composition is useful for inhibiting growth of cancer cells in an individual or for delivering a therapeutic agent to cancer cells in an individual. The invention is also useful for diagnosing cancer, for treating cancer and for inhibiting expression of a CA gene
   The invention relates to a
   Disclosure; SEQ ID NO 938; 198pp; English.
  23-SEP-2003; 2003US-00669920
  23-SEP-2004; 2004WO-US031617
  07-APR-2005
  cytostatic; gene; ds.
   Diagnosis; DNA microarray;
   16-JUN-2005
  ADZ13418;
  WO2005031001-A2
  (CHIR )
  Local Similarity
   13201 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT 13249
   3073
  2005-273395/28
  CHIRON
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT
   Conservative
   (first entry)
  DNA; 85920
  100.0%;
   1.6%;
   nucleic acid array for detecting a cancer
   microarray; biochip; cancer; neoplasm;
   0,
   Score
  Score 49;
Pred. No.
   Mismatches
  DB 4; L
   0
   Length 21477;
   Indels
   3121
   0,
   Gaps
   <u>,</u>
```

Query Match

1.68;

Score 49;

DB 12;

Length 243428;

```
RESULT 69
ADF51132/c
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   ន្តន្តន្ត
                            target. Specifically, it refers to the guanine-nucleotide exchange factor (GEF) named P-Rexl, which has also been identified as a phosphatidylinositol(3,4,5)P3-sensitive activator of Rac (a monomeric CP phosphatidylinositol(3,4,5)P3-sensitive activator of Rac (a monomeric CP phosphatidylinositol(3,4,5)P3-sensitive activator of Rac (a monomeric CP phosphatidylinositol(3)P3-sensitive activator of Rac (a monomeric CP phosphate) in returning the present invention describes this protein as CP contains in neurophils. The present invention describes this protein as CP ceducing or inhibiting inflammation, metastasis, septic shock, reducing or inhibiting inflammation, metastasis, septic shock, concluding antiinflammatory, cytostatic, an have various activities including antiinflammatory, cytostatic, antibacterial, continuosuppressive, neuroprotective and antiarteriosclerotic. Furthermore, the protein or its mutant, the nucleic acid or appropriate antibody may CP be used in a screening assay to identify a modulator of P-Rexl binding, activity or expression. This polynucleotide is the human P-Rexl genomic CP DNA sequence of the invention.
   Query Match
  human; P.Rex1; Rac; guanine-nucleotide exchange factor; GBF; GTI inflammation; metastasis; septic shock; neurodegeneration; atherosclerosis; antiinflammatory; cytostatic; antibacterial; immunosuppressive; neuroprotective; antiarteriosclerotic; gene;
Sequence
   This invention relates to a novel protein useful as an anti-inflammatory
  Disclosure;
   New isolated P-Rex1 protein or its derivative useful for discovering drugs capable of reducing or inhibiting inflammation, metastasis, sept shock, neurodegeneration or atherosclerosis, or for identifying P-Rex1
  Stephens L,
   Human P-Rex1 genomic DNA sequence
  ADF51132 standard; DNA; 243428
   21-MAR-2002; 2002GB-00006684
  21-MAR-2003; 2003WO-GB001238.
  WO2003080664-A1
   Homo sapiens
   12-FEB-2004
   ADF51132;
   (BABR-) BABRAHAM INST
   cell. This sequence represents human cancer-associated genomic DNA of
  Local
   45176
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT 3121
   invention.
  2004-011515/01.
DB; ADF51119.
  al Similarity
49; Conserv
 243428
   85920 BP; 23268 A;
   AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT
  SEQ
  Hawkins
   Conservative
   (first entry
  ij
BP; 65880 A; 63219 C; 59010 G;
  NO 14; 198pp; English.
  1.6%;
  PŢ
   0
   18962 C;
  Score 49;
Pred. No.
  ВP
   Mismatches
  19343
   3.7e-12;
  DB 14; Length 85920;
   9
  24347
 55319 T; 0 U; 0 Other;
  Indels
   7,
  0
  GEF; GTPase;
   u; o
   45128
   <u>..</u>
   ds.
   Gaps
   septic
   0
```

| NAMES/465/C  ID ANKS2458 etandard; DNA; 4316 BP.  XX  ANKS2458;  XY  O7-NOV-2001 (first entry)  XX  Human immune/haematopoietic antigen genomic sequence SEQ ID NO.37270.  XX  KY  Human immune/haematopoietic jimmune/haematopoietic antigen; cancer;  XX  VY-Costatic; gene therapy; vaccine; metastasis; ds.  XX  VY-COSTATIC; gene therapy; vaccine; metastasis; ds.  XX  XX  31-ANH-2001; 2001WO-US901354.  XX  XX  XX  31-ANH-2001; 2001WO-US901354.  XX  XX  31-ANH-2001; 2001WO-US901354.  XX  XX  31-ANH-2001; 2001WO-US901354.  XX  XX  XX  XX  XX  XX  XX  XX  XX                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | Best Local Similarity 100.0%; Pred. No. 3.6e-12; Matches 49; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  Qy 3074 GATTGTGCCACTGCAGCTTGGGCAACAGAGCAAGACTCTGTCTC 3122 |
|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| PR 12-SEP-2000; 2000US-02239979; PR 14-SEP-2000; 2000US-02239979; PR 14-SEP-2000; 2000US-02239979; PR 14-SEP-2000; 2000US-02239979; PR 14-SEP-2000; 2000US-0223998; PR 14-SEP-2000; 2000US-02330659; PR 14-SEP-2000; 2000US-02349979; PR 14-SEP-2000; 2000US-02349979; PR 14-SEP-2000; 2000US-02349979; PR 21-SEP-2000; 2000US-0236369; PR 21-SEP-2000; 2000US-0236369; PR 21-SEP-2000; 2000US-0236369; PR 22-SEP-2000; 2000US-0241809; PR 22-SEP-2000; 2000US-02446179; PR 22-SEP-2000; 2000US-0246529; PR 22-SEP-2000; | 08-SEP-2000; 2000U<br>08-SEP-2000; 2000U<br>08-SEP-2000; 2000U<br>08-SEP-2000; 2000U<br>08-SEP-2000; 2000U<br>08-SEP-2000; 2000U<br>08-SEP-2000; 2000U                  |

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AAK82461/c
ID AAK82461 standard; D
XX
AC AAK82461;
XX
DT 07-NOV-2001 (first
XX
DB Human immune/haematc
XX
KW Human; immune; haema
KW cytostatic; gene the
XX
   밁
   ঠ
  RESULT 71
   AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) CC amino acid sequences given in AAM62170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the proteins and polynucleotides may be used to proteides may be used to provent; and diagnose and treat immune/haematopoietic derived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK54942 to AAK34950 and AAM83169
  Query Match
Best Local S
Matches 48
  17.NOV-2000;
17.NOV-2000;
17.NOV-2000;
17.NOV-2000;
17.NOV-2000;
17.NOV-2000;
01.DEC-2000;
01.DEC-2000;
05.DEC-2000;
05.DEC-2000;
05.DEC-2000;
05.DEC-2000;
06.DEC-2000;
06.DEC-2000;
08.DEC-2000;
                 Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis; ds.
   Nucleic acids encoding useful for preventing,
   Sequence 4316 BP; 1270 A; 791 C; 832 G; 1423 T; 0 U; 0 Other;
  Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37273.
   Disclosure; SEQ ID NO 37270; 3071pp + Sequence Listing; English
   WPI; 2001-483426/52.
  Rosen
   (HUMA-) HUMAN GENOME SCI INC
  372 ATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGACTCTGTCTC 325
   ξ
  48;
  Similarity
   sequences
  Barash SC,
  ; 2000US-0249245P,
2000US-0249264P,
2000US-024925P,
2000US-024929P,
2000US-024929P,
2000US-0250160P,
2000US-0250160P,
2000US-0251030P,
2000US-0251989P,
2000US-0251479P,
2000US-0251868P,
2000US-0251868P,
2000US-0251868P,
2000US-0251869P,
2000US-0251989P,
2000US-0251989P,
2000US-0251999P,
2000US-0251999P,
2000US-025199P,
2000US-025199P,
2000US-025199P,
2000US-0259678P.
  ilarity 100.0%;
Conservative (
  (first entry)
  DNA;
   used in the exemplification of the
  1.5%;
   human immune/hematopoietic antigen polypeptides, diagnosing and/or treating cancers and metastasis.
   Ruben SM;
   4316
  Score 48; DB;
; Pred. No. 1.2;
0; Mismatches
  0;
  ВP
  1.2e-11;
  DB 4;
  0;
  Length 4316;
  Indels
   present invention
   0,
  Gaps
  0;
    30-JUN-2000

07-JUL-2000

11-JUL-2000

11-JUL-2000

12-JUL-2000

12-JUL-2000

14-JUL-2000

14-JUL-2000

14-JUG-2000

16-JUG-2000

17-JUG-2000

18-JUG-2000

18-JUG-2000

18-JUG-2000

18-JUG-2000

19-JUG-2000

10-SEP-2000

10-SEP-2000

10-SEP-2000

10-SEP-2000

10-SEP-2000

10-SEP-2000

11-SEP-2000

21-SEP-2000

   11-JAN-2000
04-FEB-2000
24-FEB-2000
02-MAR-2000
16-MAR-2000
17-MAR-2000
18-APR-2000
19-MAY-2000
   07-JUN-2000;
28-JUN-2000;
  17-JAN-2001;
   Homo sapiens.
   WO200157182-A2
 2000US-0186528P
2000US-0186350P
2000US-0186350P
2000US-019813P
2000US-0198129P
2000US-0215135P
2000US-021518880P
2000US-021518880P
2000US-0217487P
2000US-0224518P
2000US-0225513P
2000US-0225513P
2000US-0225513P
2000US-0225513P
2000US-0225513P
2000US-0225513P
2000US-0225513P
2000US-0225513P
2000US-0225758P
2000US-0225758P
2000US-0225758P
2000US-0225758P
2000US-0225758P
2000US-0225758P
2000US-0225759P
2000US-0225739343P
2000US-023343P
2000US-023343P
2000US-02331343P
2000US-02331413P
2000US-02331439P
2000US-023342349P
2000US-023342349P
2000US-023342349P
2000US-0233499P
2000US-0233499P
2000US-0233499P
  2001WO-US001354
   2000US-0179065P
```

2000US-0235834P. 2000US-0235836P. 2000US-023637P. 2000US-0236367P. 2000US-0236368P. 2000US-0236369P.

```
RESULT 72
AAK82456/c
  CC AAKS4951 to AAK64702 encode the human immune/haematopoietic antigen (I) camino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic CC activity, and can be used in gene therapy and vaccine production. (I) crosselins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I) by inserting the CC nucleic acids into a host cell and culturing the cell to express the CC diagnose and treat immune/haematopoietic-derived cells. AAK64703 cancers and cancer metastases of haematopoietic antigen genomic sequences from the present invention. AAK54942 to AAK87654 and AAM82169 creatment served in the arreant invention. AAK54942 to AAK8765 and AAM82169
   밁
   á
  Query Match
Best Local S
Matches 48
31-JAN-2000

04-FEB-2000

24-FEB-2000

24-FEB-2000

12-MAR-2000

16-MAR-2000

17-MAR-2000

19-MAY-2000

07-JUN-2000

28-JUN-2000

07-JUL-2000

07-JUL-2000
  Sequence 4316 BP; 1270 A; 791 C; 832 G;
   Disclosure; SEQ ID NO 37273; 3071pp + Sequence Listing; English
  Nucleic acids encoding
  07-NOV-2001
   AAK82456;
   Rosen
   09-AUG-2001
   Human immune/haematopoietic antigen genomic sequence
  AAK82456
   represent sequences used in the exemplification of the
  17-JAN-2001;
  WO200157182-A2
  cytostatic;
  Human; immune; haematopoietic; immune/haematopoietic antigen; cancer
  Local Similarity
   3075 ATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  2001-483426/52
   372
  Ç
   48;
  for
  standard;
   ATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  preventing,
   Barash SC,
  1.5%; Scilarity 100.0%; P
Conservative 0;
2000US-018464P.
2000US-0189874P.
2000US-0199076P.
2000US-019123P.
2000US-0205515P.
2000US-0205467P.
2000US-02154866P.
2000US-0215435P.
2000US-0216647P.
2000US-0216680P.
  gene
  2000US-0179065P.
2000US-0180628P.
   2001WO-US001354
  (first entry)
  therapy; vaccine; metastasis; ds.
  DNA;
  human immune/hematopoietic antigen polypeptides, diagnosing and/or treating cancers and metastasis.
   Ruben
  4317
   Score 48;
Pred. No.
   SM:
  ₽₽
   Mismatches
  1423 T; 0 U; 0 Other;
  DB.
   .2e-11;
  ..
   Length 4316;
   Indels
  SEQ ID NO: 37268
   present
   325
   0
  invention
   Gaps
```

0

(HUMA-)

NAMOH

GENOME

SCI

27-SEP-2000;
27-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
29-SEP-2000;
20-CT-2000;
20-NOV-2000;
20-NOV-2000;
20-NOV-2000;
20-NOV-2000;
20-NOV-2000;
20-NOV-2000;
20-NOV-2000;
20-NOV-2000;
20-NOV-2000;
21-NOV-2000;

2000US-0236370P.
2000US-023703P.
2000US-023703P.
2000US-023703P.
2000US-023703P.
2000US-023703P.
2000US-023703P.
2000US-0241966P.
2000US-0241808P.
2000US-0246478P.
2000US-0246478P.
2000US-0246478P.
2000US-0246478P.
2000US-0246523P.
2000US-0246523P.
2000US-0246523P.
2000US-0246523P.
2000US-024651P.
2000US-024921P.
2000US-025198P.
2000US-0251186P.
2000US-0251186P.
2000US-0251186P.
2000US-0251186P.
2000US-0251186P.
2000US-0251186P.
2000US-0251186P.
2000US-0251189P.
2000US-0251198P.
2000US-0251186P.
2000US-0251198P.
2000US-0251198P.
2000US-0251186P.
2000US-025186P.
2000US-025186P.
2000US-025186P.
2000US-025186P.
2000US-025186P.
2000US-025186P.
2000US-025186P.
2000US-025186P.

```
20-OCT-2000
01-NOV-2000
08-NOV-2000
17-NOV-2000
17-NOV
AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis are treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the
   Disclosure;
  (HUMA-)
  2001-483426/52
   Š
  c acids encoding for preventing,
  HUMAN
   Barash
   SEQ.
   2000US-0241826P

2000US-024647PP

2000US-024647PP

2000US-024647PP

2000US-024647PP

2000US-0246524P

2000US-0246525P

2000US-0246528P

2000US-024652PP

2000US-024652PP

2000US-024651PP

2000US-024661PP

2000US-024661PP

2000US-024661PP

2000US-024921PP

2000US-0251988PP

2000US-02511869P

2000US-02511869P

2000US-0251989P

2000US-0251989P

2000US-025199PP

2000US-025199PP

2000US-025199PP

2000US-025199PP

2000US-025199PP

2000US-025199PP
  GENOME
   Ħ
   SC,
   ĕ
  SCI
   37268;
  human immune/hematopoietic diagnosing and/or treating
   Ruben
   3071pp +
   NS
   Sequence
   Listing;
  antigen
cancers
```

English

and

the

polypeptides, and metastasis.

11-JUI-2000 11-JUI-2000 12-JUI-2000 12-JUI-2000 11-JUI-2000 11-JU

2000US-0217487P
2000US-021964P
2000US-022964P
2000US-0225213P
2000US-0225213P
2000US-0225213P
2000US-022526P
2000US-022526P
2000US-0225276P
2000US-0225276P
2000US-0225759P
2000US-0225759P
2000US-02257686P
2000US-0225759P
2000US-0231244P
2000US-0231244P
2000US-0231244P
2000US-0231244P
2000US-0231244P
2000US-0231244P
2000US-0231244P
2000US-0231244P
2000US-02312499P
2000US-0235836P
2000US-0235836P
2000US-0235836P
2000US-0235839P

ពិនិត្តនិត្តនុំនុំ

밁 S

```
The invention relates to recombinant carcinoma associated (CA) nucleic CC acid sequences from mouse and human (AAA01492-ADA03094), and to CC acid sequences from mouse and human (AAA01492-ADA03094), and to CC combinant carcinoma associated proteins (CAP) encoded by them. The CC invention also encompasses expression vectors and host cells comprising a CC canucleic acid, a polypeptide (especially an antibody) that specifically binds to the protein, and a blochip comprising CA nucleic acid or concequents thereof. The sequences of the invention were identified using CC oncogenic retroviruses, which insert into the genome of the host organism CC at random. Many of these do not carry transduced host oncogenes or pathogenic trans-acting viral genes, meaning that cancer incidence is a CC direct consequence of the effects of proviral integration into host CC protooncogenes. The CA nucleic acid sequences can be used to diagnose CC carcinoma (especially breast cancer, prostate cancer, lymphoma or carcinoma (especially breast cancer, prostate cancer, lymphoma or complexements) or a propensity to carcinoma by determination of the sequence of a cacids, proteins and antibodies are also useful as the transcutic agents and in screening and evaluating drug candidates. The presents sequence of the invention. Note: The complete sequence data for this gatent did not form part of the printed specification, but was obtained
   Matches
  Query Match
Best Local :
  protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-derived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK54942 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention
   Sequence 4317 BP; 1270 A; 791 C; 831 G; 1425 T; 0 U; 0 Other;
   17-JUL-2003.
   Claim 1; SEQ ID NO 1184; 245pp; English.
  New recombinant nucleic acid encoding carcinoma associated protein, useful for preparing compositions for treating carcinomas.
  Morris
   (SAGR-) SAGRES DISCOVERY
  26-DEC-2001; 2001US-00035832
  26-DEC-2002; 2002WO-US041414.
  WO2003057146-A2
   Homo sapiens
  prostate; lymphoma;
  Human; carcinoma associated; oncogene; carcinoma; cancer; breast;
   Human MDM2 carcinoma associated gene,
  06-NOV-2003
   ADA02666;
  ADA02666 standard; DNA; 52242
  Local
  3075 ATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   2003-587068/55
  372
   48;
  Similarity
  ATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 325
   1.5%;
larity 100.0%;
Conservative (
  (first
  leukaemia; cytostatic; gene therapy; drug screening;
  entry
   score 48; DB; Pred. No. 1.2
  ВP
   SEQ ID NO:1184
  DB 4;
  ..2e-11;
   0,
  Length 4317;
   Indels
   0
   Gaps
   0
```

RESULT 75 ADE95914

```
ADB72404
ADB72X0
ADB72X0
AC ADB7
AC ADB7
AX AD
  밁
  S
   នន្តន្ត
밁
  5
  Query Match
Best Local Similarity
   Matches
   Matches
   Query Match
Best Local Similarity
  The invention relates to a novel recombinant nucleic acid comprising a nucleotide sequence selected from any of the 660 sequences fully defined in the specification. A polynucleotide of the invention has cytostatic activity, and may have a use in gene therapy, or in a vaccine. The recombinant nucleic acids and polypeptides are useful for treating carcinomas, e.g. lymphomas, cancers, neoplasm, adenocarcinoma, and
   02-MAR-2001; 2001US-00798586.
23-CCT-2001; 2001US-00004113.
08-NOV-2001; 2001US-00052482.
30-NOV-2001; 2001US-00997722.
20-DEC-2001; 2001US-00034650.
  ADB72404;
  in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
   30-JAN-2003
  Homo sapiens
  cancer; neoplasm;
   human; ds; cytostatic; gene therapy; vaccine; carcinoma; lymphomas; cancer; neoplasm; adenocarcinoma; sarcoma; gene.
  Human MDM2 gene
  04-DEC-2003
   ADB72404
  Sequence 52242 BP; 14384 A; 10354 C; 10997 G; 16487 T; 0 U; 20
   Claim 1;
  cancers,
  WPI; 2003-239337/23
   sarcomas.
  New recombinant nucleic acid, useful for treating carcinomas, lymphomas,
   26-DEC-2001; 2001WO-US051291
  WO2003008583-A2
   (SAGR-) SAGRES DISCOVERY
  5520
  5520
  2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
   48;
   W.
   48;
GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
                             GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
   SEQ ID NO 232; 2304pp; English.
  standard;
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
   52242 BP; 14384 A; 10354 C; 10997 G; 16487 T; 0 U;
   neoplasm,
   The present sequence represents a human gene
   Engelhard
  Conservative
   Conservative
  (first
   adenocarcinoma,
  DNA;
   1.5%;
  100.0%;
  1.5%;
  묫
   52242
   0
   0
  Score 48; DB 10;
Pred. No. 1.1e-1;
   Score 48;
Pred. No.
  ВP
   Mismatches
  Mismatches
  or sarcomas
  DB 9;
  1.1e-11;
   .1e-11;
  Length 52242;
  Length 52242;
   Indels
  Indels
  ę,
  2942
  2942
   5567
  5567
  0,
  20 Other;
   0
  invention
   Other;
   Gaps
  Gaps
   0
  0
```

```
RESULT 76
AEAG1175/c
ID AEAG11
XX AEAG11
XX AEAG11
XX AEAG11
XX DEAG11
XX DEAG11
XX DEAG11
XX DEAG11
XX Human
XX Human
XX DNA me
XX DNA me
XX DNA me
XX US2005
XX US2005
XX JONO 8
   S
  밁
   Matches
   Query Match
Best Local Similarity
  diagnosis and treatment of cancer, especially carcinomas, as well as the use of compositions in screening methods. The compositions of the invention may have cytostatic activity whilst the disclosed sequences may be useful for gene therapy. The carcinoma associated nucleic acids and proteins are useful for diagnosing and treating carcinomas, for example lymphoma, breast cancer, prostate cancer or leukaemia, or for screening drug candidates or bioactive agents capable of binding to, or modulating the activity of, a carcinoma associated protein. The present sequence is the genomic DNA sequence of the human MDM2 gene which is a carcinoma
   Human ENTPD5 gene genomic sequence SEQ ID NO:85.
   Sequence 52242 BP;
  Claim 1; SEQ ID NO 172; 793pp; English.
  New carcinoma associated nucleic acids and proteins, useful for screening drug candidates, or for diagnosing and treating carcinomas, e.g. lymphoma, breast cancer, prostate cancer or leukemia.
   Morris DW,
  08-NOV-2001; 2001US-00052482
   08-NOV-2002; 2002WO-US036071.
  WO2003039484-A2
   Homo sapiens
  cancer diagnosis;
   Human MDM2 gene genomic DNA sequence
   16-JUN-2005
  25-AUG-2005
  ABA61175
  AEA61175 standard; DNA; 53779 BP
  associated gene of the invention.
   WPI; 2003-441462/41
  15-MAY-2003
  ADE95914 standard; DNA; 52242
   Homo sapiens
  DNA methylation;
   This invention relates to novel recombinant nucleic acids for use
  (SAGR-) SAGRES DISCOVERY.
   lymphoma;
   2895
   48;
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 5567
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
   breast
  Conservative 0;
   Engelhard
  (first entry)
  (first entry)
  ; cancer treatment; carcinoma; cytostatic; gene therapy; cancer; prostate cancer; leukaemia; ds; human; MDM2.
  biomarker; cancer; gene; ds; ENTPD5
   14384 A; 10353 C; 10998 G; 16487 T; 0 U; 20 Other;
   1.5%;
   묫
   Score 48;
Pred. No.
   Mismatches
  DB 10; I
1.1e-11;
hes 0;
   Length 52242;
  Indels
   2942
  0
  Gaps
   Ħ
  <u>.</u>
```

WO2003073826-A2 Homo sapiens Cytostatic; carcinoma; lymphoma; cancer;

human;

0

Human genomic sequence hCG16651.

```
RESULT 77
ACN44374
ID ACN44374
XX ACN44
XX ACN44
XX Humal
XX Humal
XX Cytor
XX Cytor
XX Homo
X
   片
   S
   CC methylation of one or more CpG sites on nucleic acid sequences in a CC biological sample obtained from the subject, and determining the presence CC of, predisposition to, or stage of the disease in the subject based on CC the degree of methylation; (2) monitoring the onset, progression, or CC regression of a disease in a subject; (3) determining the efficacy of a clest compound for inhibiting a disease in a subject; and (4) a kit (1) CC useful for diagnosis, prognosis, staging, monitoring, and therapeutic CC treatment of a disease. (M1) is useful for identifying one or more CC nucleic acid sequences useful as a biomarker for a disease to be CC detected, where the nucleic acid sequences are useful for detecting, the present sequence represents a specifically claimed human CC genomic sequence for use in the method of the invention. Note - The sequence data for this patent is not represented in the printed CC specification but was obtained in electronic format from the USPTO web
   Query Match
Best Local Similarity
Matches 48; Conserv
   nucleic acid sequences useful as a biomarker for a disease to be detected. (M1) involves identifying nucleic acid sequences comprising methylated CpG site in promoter-first exon region and that are down-regulated in diseased cells, comparing expression level of nucleic acid sequences with that of demethylated nucleic acid sequences and identifying nucleic acid sequences exhibiting increase in expression after demethylation. Also described: (1) detecting (M2) the presence or stage of a disease in a subject, which involves determining the degree of the described of the comparison of 
  Sequence 53779 BP; 14286 A; 11767 C; 12248 G; 15478 T; 0 U; 0 Other;
   Claim 11; SEQ ID NO 85; 27pp; English.
   Identifying nucleic acid sequences as biomarker for disease, by identifying nucleic acid sequences comprising methylated CpG site and down-regulated in diseased cells and comparing its expression level with demethylated nucleic acid.
   18-NOV-2004
   ACN44374;
   ACN44374 standard; DNA; 181684 BP
   The invention
  GENBANK; NM_001249.
   WPI; 2005-456991/46
  Beard C,
   16-DEC-2003; 2003US-00737082
   27-JAN-2004; 2004US-00765790
   2311 GAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCA
   GAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCA
   Burgess C,
   1.5%; So ilarity 100.0%; I Conservative 0;
   (first entry)
   relates to a method (M1) for identifying one or
   CORP.
   Gannon A,
  Score 48;
Pred. No.
   Mismatches
   Harvey J,
  DB 14; I
. 1.1e-11;
   Lechner JF,
  Length 53779;
  Indels
  드
   2264
  2936
   0
   2
   Gaps
```

of.

```
XGXGXGXGXGXGX
  밁
  á
   Query Match
Best Local S
Matches 48
   16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
  The present invention relates to novel DNA and protein sequences which are associated with carcinomas. The sequences are useful for: (1) for screening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (1ii) for screening of a bioactive agent capable of modulating the activity of CAP; (iv) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing carcinoma; (vi) for inhibiting the activity of CAP; (ix) as a biochip; (x) for diagnosing carcinoma; (vii) for neutralizing the effect of CAP; (1x) as a biochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the
  Recombinant nucleic acid useful for diagnosis and treatment of carcinoma comprises a nucleotide sequence.  \begin{tabular}{ll} \hline \end{tabular} 
   uscermining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent US200218258AA1 for which no common the patent to basic patent
  17-FEB-2000;
  13-SEP-2002
  Sequence 181684 BP; 55185 A; 34753 C; 35001 G; 55847 T; 0 U; 898 Other;
  Claim 1; SEQ ID NO 790; Opp; English.
   WPI; 2003-328604/31
  28-FEB-2003;
                                 (MILL-)
  20-FEB-2001; 2001WO-US005171.
  WO200160860-A2
   Human; prostate cancer; cytostatic; carcinogen; pharmacodyanamic marker; pharmacogenomic marker; gene; ss.
  Human prostate expression marker cDNA 16322.
  ABV16331;
   ABV16331 standard; cDNA;
  US2002182586A1,
   (SAGR-)
  155726
   3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTC 3120
  sapiens
   48;
   SAGRES
                                MILLENNIUM PREDICTIVE MEDICINE INC.
   Similarity
×
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTC 155773
  ; 2000US-0183319P.
; 2000US-0189862P.
; 2000US-0207454P.
; 2000US-0211314P.
; 2000US-0219007P.
; 2000US-0255281P.
   Conservative
   (first
  2002US-00087192
  2003WO-US006235
   DISCOVERY
   for which no sequence data was published
Š
   entry
  1.5%; Score 48; DB
100.0%; Pred. No. 1.:
tive 0; Mismatches
   440
Monahan JE
   ВP
   DB 11; I
1.1e-11;
   0;
  Length 181684;
   Indels
  0,
   Gaps
   0
```

```
RESULT 79
ABV46129
ID ABV46
XX ABV46
XX ABV46
XX I-SE
DT 16-SE
XX Human
XX Phane
XX I-FE
XX 1-FE
XX 1-FE
XX 1-FE
XX 1-FE
XX 1-FE
XX 11-FE
PR 13-DE
PR 13-DE
PR 13-DE
PR 13-DE
PR 13-DE
XX WPI;
XX WOYel
PT POSET
XX NOVel
PT POSET
XX Claim
   XXCCCCCCCCCCCXXXXTTTXXX
  밁
   S
  Best Local Similarity Matches 47; Conserv
  Query Match
   16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
  a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a patient; (l) is also useful as a pharmacodyanamic or pharmacogenomic marker
   Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer.
   Human; prostate pharmacogenomic
  Human
  16-SEP-2002
   ABV46129
   Sequence
   Claim 1; Page 2728; 11750pp; English
   Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful
  WPI; 2001-662795/76.
   Homo sapiens.
  specification or
   The invention relates to
   18-JUL-2000;
13-DEC-2000;
  17-FEB-2000;
  20-FEB-2001; 2001WO-US005171
   23-AUG-2001.
   WO200160860-A2
   (MILL-) MILLENNIUM PREDICTIVE MEDICINE
  nucleotide sequence given in Tables
   detecting
   detecting
  3070 GCAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
   prostate expression marker cDNA 46120
  æ
   standard; cDNA; 516
   440 BP; 136 A; 96 C; 112 G; 96 T; 0 U; 0 Other;
  ; 2000US-0183319P.
; 2000US-0189862P.
; 2000US-0207454P.
; 2000US-0211314P.
; 2000US-0219007P.
; 2000US-0255281P.
  ilarity 100.0%;
Conservative
  (first entry)
  Endege
   presence of prostate cancer, stage of prostate
   presence
   cancer; cytostatic; carcinogen; pharmacodyanamic marker;
marker; gene; ss.
  its complement. (I) is useful for: (a) assessing whether
  ģ
   1.5%;
   an isolated nucleic acid molecule (I) comprising
  0;
  Score 47;
Pred. No.
   ВÞ
  Mismatches
  1-9 (ABV00010-ABV62213) of the
  DB 5; Le
  Length 440;
  Indels
  0
  Gaps
  0
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9110; 11750pp; English

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RESULT 80
ADL13941
ID ADL13
XX ADL13
AC ADL13
XX OSTEO
XX
   Best Loca
Matches
The invention relates to a method of determining susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain comprising identifying whether the individual has at least one polymorphism in a polymucleotide encoding at least one of the protein listed in the specification. The methods, composition and agent are useful for modulating the susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain that is associated with a disease, preferably osteoarthritis. The cell line and the non-human animal are useful for screening for an agent for diagnosing an individual having susceptibility to joint space narrowing and/or osteophyte development and/or joint space narrowing and/or to the polynucleotide encoding a protein listed in the specification. (Note:
   The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for: (a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a patient assessing the aggressiveness or indolence of prostate cancer in a patient.
   Disclosure; SEQ ID NO 473; 297pp; English.
  Determining susceptibility of an individual to joint space narrowing, osteophyte development and/or joint pain comprises identifying whether the individual has at least one polymorphism in a polynucleotide encoding
   Jones
   20-DEC-2001; 2001US-0342603P
   19-DEC-2002;
   03-JUL-2003.
   WO2003054166-A2
   Homo sapiens
  Osteoarthritis-associated polymorphic nucleotide
  06-MAY-2004
  ADL13941 standard; DNA; 125515
  Sequence 516 BP; 161 A; 118 C; 135 G; 100 T; 0 U; 2 Other;
   (INCY-)
  ADL13941;
  Local
   Match
  gene; osteopathic; antiinflammatory; antiarthritic; gene therapy;
   3070
  2003-559141/52.
   418
   Š
   18
   space narrowing; osteophyte development; joint pain;
   Similarity 100.
47; Conservative
   INCYTE
   GCAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 464
   GCAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  also
   Schafer A;
   2002WO-US041225.
  (first entry)
  useful as a pharmacodyanamic or pharmacogenomic marker
   GENOMICS
   SNP; single
  100.0%;
   1.5%;
   nucleotide polymorphism.
   .
.
  Score 47;
Pred. No.
  ВP
   Mismatches
  DB 5; Lo
  Length 516;
   Indels
   0
   diagnosing
   Gaps
  encoding
   one
   0
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ARBSULT 81
ADW6605/c
ID ADW660
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  8
  នួននេះ
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   Query Match
Best Local
  schizophrenia; psychosis; alopecia; inflammatory disorder; attaxia telangiectasia; diabetes; skin disorder; osteoarthritis; rheumatoid arthritis; blood pressure; atherosclerosis; cardiovascular disease; pulmonary disease; Alzheimers disease; parkinsons disease; osteoporosis; asthma; developmental disorder; infertility; infection; cytostatic; immunosuppressive; dermatological; antiinflammatory; neuroprotective; gastrointestinal-gen; neuroleptic endocrine-gen; antidiabetic; antiarthritic; osteopathic; antirheumatic; antiarteriosclerotic; cardiovascular-gen; nootropic;
   The sequence data for this patent did not form specification but was obtained in electronic for ftp.wipo.int/pub/published_pct_sequences).
   New human gene trapped sequences, useful for diagnosing and treating disorders affecting development and cell differentiation, e.g. aging cancer, schizophrenia, alopecia, diabetes, rheumatoid arthritis, or
  ADW06065 standard; DNA;
   Nehls M,
  30-0CT-1998;
27-0CT-1999;
   06-AUG-2004; 2004US-00914037
   06-JAN-2005
  US2005003444-A1
   systemic lupus erythematosus; scleroderma; crohns disease; multiple sclerosis; inflammatory bowel disease; immune disorder;
  gene expression;
systemic lupus e
  Human gene trapped sequence (GTS)
   (LEXI-) LEXICON GENETICS INC
  antiparkinsonian;
   3076
  2005-065239/07
   sapiens.
   l Similarity
47; Conserv
  TTGTGCCACTGCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   TTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   125515 BP;
  Zambrowicz
  Conservative
  (first entry)
  98US-0106442P.
99US-00428674.
  forensic; aging; cancer;
  antiasthmatic;
   33180 A;
  100.0%;
  ₽,
  1.5%;
   380
  0,
  Score 47;
Pred. No.
   28822 C;
  Ą
  Mismatches
  antiinfertility;
   - SEQ ID
   28744 G;
  DB 10;
  3.2e-11;
  autoimmune disease;
  282.
  crohns disease;
  cm part of the printed format directly from WIPO at
   34769 T;
   Length 125515;
  gene trapped sequence;
  Indels
  0 U; 0 Other;
   3122
  77339
  0
  Gaps
```

The invention comprises novel human gene trapped sequences (GTSs) which are useful in gene discovery and as markers for gene expression analysis, forensic analysis, and determining the genetic basis of human disease. The human GTSs of the invention are useful for diagnosing and treating disorders affecting development and cell differentiation, such as: aging, cancer, autoimmune disease, hupus, scleroderma, Crohn's disease, multiple sclerosis, inflammatory bowel disease, immune disorders, schizophrenia, psychosis, alopecia, glandular disorders, inflammatory disorders, ataxia telangiectasia, diabetes, skin disorders, osteoarthritis, rheumatoid

Claim 3;

SEQ

ID NO 282;

35pp; English

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RESULT 82
ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABL83399/c
ID ABCBC
ID 
cc and immunostitudiate; and a polypeptide (II) of a ovarian tumour (S1) colleges of the 10912 nucleotide sequences as given in ABL77023 to ABL87934, CC (III) encoding (II) having a sequence (S2), a T cell population of (II), CC or antigen presenting cells that express (II). (I) has cytostatic contribute. An oligonucleotide (IV) that hybridises to (S1) can be used for detecting ovarian cancer in a patient's biological sample preferably comparing the amount of polynucleotide (IV), detecting the amount of polynucleotide (IV), detecting the amount of polynucleotide comparing the amount to a predetermined cutoff comparise of (IV) and comparing the amount to a predetermined cutoff composition of polynucleotide hybridising to (IV) is detected preferably by colleges of polynucleotide hybridising to (IV) is detected preferably by colleges of polynucleotide hybridising to (IV) is detected preferably by colleges of polynucleotide hybridising to (IV) is detected preferably by colleges of polynucleotide hybridising to (IV) is detected preferably by college of polynucleotide hybridising to (IV) is detected preferably by colleges of polynucleotide profess of the tumour colleges of the tumour colleges of the tumour colleges of the tumour polypeptides and protesins in tumour cells; and colleges of the tumour colleges and protesins in tumour cells; and colleges of the tumour colleges of the colleges of the tumour colleges of the co
   Query Match
Best Local S
Matches 46
   arthritis, high blood pressure, atherosclerosis, cardiovascular disease, pulmonary disease, degenerative disease of neural or skeletal systems, Alzheimer's disease, Parkinson's disease, osteoporosis, asthma, developmental disorders, genetic birth defects, infertility, epithelial ulcerations, and infections. The present nucleic acid represents a human GTS of the invention. NOTE: The present sequence is not shown in the specification, but has been retrieved from the USPTO web site.
   polypeptide of a ovari polypeptide, antibody polypeptide.
   Composition
  Algate
  29-MAY-2001; 2001WO-US017756
   06-DEC-2001.
  Human ovarian cancer related cDNA clone SEQ ID NO:6376
  Sequence 380
  Claim 1; SEQ ID NO 6376; 489pp; English
   WPI; 2002-122075/16.
  26-MAY-2000; 2000US-0207484P
  WO200192581-A2
   Homo sapiens
   Human; ovarian
   17-MAY-2002
  ABL83398;
  ABL83398 standard; cDNA; 405
   (CORI-) CORIXA CORP
  present invention immunostimulants;
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
   49
   1 Similarity
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
  Harlocker
  of a ovarian tumor, antibody specific
   Conservative
   for therapy and diagnosis of ovarian cancer comprising
  вр;
   (first entry
  cancer; ovarian tumour; cytostatic; gene; ss
  93 A; 103
   100.0%; ---
   describes a composition
  SL,
  Ç,
  Jones
   Score 46; DB; Pred. No. 1.10; Mismatches
  쁑
  78 G;
  polypeptide, polynucleotide encoding to polypeptide or T cell expressing
  ₽,
  105 T; 0 U; 1 Other;
   1.1e-10;
hes 0;
  14;
   Ξ
  Length 380
   comprising: carriers
   Indels
   0,
   Gaps
   0
```

```
RESULT 83
ADL43370/c
ID 43370/c
ID 43370/c
ID 43370/c
ID 20-MAY
XX Human
XX Human
XX Human;
XX Human;
XX Homo 8
XX Homo 8
XX Human;
XX Homo 8
XX I - MAH
PR 21-MAH
PR 25-JUI
PR 21-JUI
PR 21-MAH
PR 
   문
  S
   SXS
   Query Match
Best Local
  Matches
                      patient afflicted with ovarian cancer comprising providing to cells of the patient an antisense oligonucleotide complementary to a marker of the invention. The markers are useful for assessing if a patient is afflicted with ovarian cancer, which involves comparing the level of expression of a marker in a patient sample and a normal level of expression of the marker in a control non-ovarian cancer sample. A difference between the expression levels indicates ovarian cancer. The level of expression of a marker corresponds to a secreted protein or to a transcribed polynucleotide or its portion. The level of expression of the marker is assessed by detecting the presence in the sample, a protein or protein fragment corresponding to the marker. The presence of protein or protein fragment is detected using an antibody that specifically binds with the protein or protein fragment is detected using an antibody that specifically binds with the protein or protein. Alternatively, the level of expression of
  21-MAR-2000;
25-MAY-2000;
15-JUN-2000;
   The invention relates to nucleic acid markers which are overexpressed in ovarian cancer cells as compared to their expression in normal (i.e. non-cancerous) ovarian cells. The invention also relates to polypeptides encoded by the markers, antibodies that selectively bind to the polypeptides, a method of inhibiting ovarian cancer in a patient at risk of developing ovarian cancer involving inhibiting expression of a gene corresponding to a marker of the invention and a method of treating a corresponding to a marker of the invention and a method of treating a
   Novel isolated nucleic acid molecules (markers) overexpressed in ovarian cancer cells as compared to their normal non-cancerous ovarian cells are used to characterize stage, grade, histological type of ovarian cancer.
  07-JUL-2000;
25-JUL-2000;
  21-MAR-2001; 2001WO-US009126
  Human ovarian cancer DNA marker #17260.
   20-MAY-2004
  ADL43370;
   ADL43370 standard; DNA; 458
  library using well known techniques
   Disclosure; SEQ ID
   WO200170979-A2
  Homo sapiens.
   (MILL-)
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGACACAGACTC 3116
   106 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 61
   46;
  ovarian
   MILLENNIUM
  Similarity
   405 BP; 84
  2000US-0207124P.
2000US-0211940P.
2000US-0216820P.
2000US-0220661P.
   Conservative
   2000US-0191031P.
   (first
  2000US-0257672P.
  cancer;
   NO 17260; 106pp; English.
  PREDICTIVE MEDICINE INC.
  entry)
   A; 114 C;
  1.5%;
      λĄ
  ds; tumour; cytostatic;
  0
  ВP
  Score 46;
Pred. No.
   95 G; 112
   Mismatches
  1.1
   T; 0
  .1e-10;
   U; O
  DNA marker
  Length 405
  Indels
   Other;
  0
  Gaps
  0
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RESULT 84
ABV60535
ID ABV60
  Query Match
Best Local
  17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for: (a) assessing whether a patient is affilicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cancer has metastatized in a patient; (f) assessing the aggressiveness or indolence of prostate cancer in a patient; (h)
  polynucleotide which anneals with the marker or anneals with a portion of the polynucleotide comprising the marker, under stringent conditions. The marker is also used for monitoring the progression of ovarian cancer in a patient which involves detecting expression of the marker in a patient time and comparing the level of expression. The method at a subsequent using an ovarian tissue sample. A composition comprising a marker, polypeptide or antibody of the invention is used to treat ovarian cancer. This sequence represents a human ovarian cancer DNA marker of the
  Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usef for detecting presence of prostate cancer, stage of prostate cancer.
   The invention relates to
  Claim 1;
   Schlegel
  20-FEB-2001; 2001WO-US005171
  23-AUG-2001
   WO200160860-A2
   Homo sapiens.
  Human; prostate pharmacogenomic
  Human prostate
   Sequence
  13-SEP-2002
   ABV60535;
   ABV60535 standard;
  (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
  Local Similarity
  2895
  405
   ₹
  Page 11527; 11750pp; English.
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 360
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
  458 BP;
   ; 2000US-0183319P.
; 2000US-0189862P.
2000US-0207454P.
; 2000US-0211314P.
; 2000US-0219007P.
; 2000US-0255281P.
   Conservative
   Endege
   (first entry)
  cancer;
  expression marker cDNA 60526.
   89
   CDNA; 497
  100.0%;
   ĕ,
  A; 124 C; 126
  1.5%;
  gene;
   cytostatic; carcinogen; pharmacodyanamic marker;
  an isolated nucleic acid molecule (I)
  Monahan JE
   0,
   ВP
  Score 46;
Pred. No.
   Mismatches
  G; 119
  1.1e-10;
  DB 5;
  T; 0 U; 0 Other;
  stage
   0;
   Length 458
   Indels
  0;
   comprising
  Gaps
   0
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RESULT 85
AAH18284/c
밁
  S
   ន្តដ្ឋន
  Query Match
Best Local
   Matches
  27-AUG-1999;
11-JAN-2000;
02-MAY-2000;
09-JUN-2000;
          oligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprising a sequence complementary to the complementary strand of a polynucleotide which comprises a 5'-end sequence and an oligonucleotide comprising a sequence complementary to a polynucleotide which comprises a 3'-end sequence, where the oligonucleotide which comprises a 1'-end sequence, where the specification. The primer sets can be used in antisense therapy and in gene therapy. The primers are useful for synthesising polynucleotides, particularly full length cDNAs. The primers are also useful for the detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs. The primers allow obtaining of the full-length cDNAs assily without any specialised methods. AAH03166 to AAH13628 and AAH13633 to AAH18742 represent human cDNA sequences; AAB92446 to AAB95893 represent human amino acid sequences. and AAH13639 to AAH13639 represent
   The present invention describes primer sets for synthesising 5602 full-length cDNAs defined in the speciation. Where a primer set comprises:

(a) an oligo-dT primer and an oligonucleotide complementary to the complementary strand of a polynucleotide which comprises one of the 5602 nucleotide sequences defined in the specification, where the
   Primer sets for synthesizing polynucleotides, particularly the 5602 length cDNAs defined in the specification, and for the detection and diagnosis of the abnormality of the proteins encoded by the full-length control of the proteins encoded by the proteins encoded by the full-length control of the proteins encoded by the full-length control of the proteins encoded by the p
  Claim
  Ota T,
   29-JUL-1999;
  28-JUL-2000;
   07-FEB-2001
  EP1074617-A2.
   26-JUN-2001
  AAH18284;
   AAH18284
   Sequence
  WPI; 2001-318749/34.
   Homo sapiens.
  Human; primer;
   Human cDNA sequence SEQ ID NO:18263
   (HELI-)
  Local
  Ξ
   2895
  192
  18
  1 Similarity
46; Conser
  Isogai T,
   HELIX RES INST.
  SEQ
   497
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
   standard;
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 237
   also useful as a pharmacodyanamic or pharmacogenomic marker
   Sugiyama T,
  99JP-00300253.
2000JP-00118776.
2000JP-00183767.
2000JP-00241899.
  日
  2000EP-00116126
  BP; 145 A; 101 C; 133 G; 118 T; 0 U; 0 Other;
  (first
  detection; diagnosis;
   99JP-00248036
  NO 18263;
  1.5%; Sur
, 100.0%; Pr
   Nishikawa T,
T, Wakamatsu
   CDNA;
   entry)
   2537pp +
  Score 46;
Pred. No.
   Mismatches
  ۶
  Hayashi K,
  Sequence
  Nagai K,
  antisense therapy; gene therapy; ss.
  DB 5; Le
1.1e-10;
sequences; in AAH13629 to
   Listing; English
  Saito K,
  Otauki
   Length 497;
   Indels
  H
  Yamamoto
   ction and/or full-length
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  Gaps
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ADV15647
ADV15647
ID 1547
AC ADV15
XX ADV156
XX ADV15
XX ADV15
XX ADV15
XX ADV15
XX ADV15
XX ADV16
XX AV16
XX AV
RESULT 87
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ID ACC82887 standard; DNA; 7001 BP.
  밁
   ঠ
   Query Match
Best Local S
Matches 46
   Query Match
  The invention relates to an isolated nucleic acid encoding a PRO polypeptide. The polypeptide, agonist or an antagonist, antibody, composition, and method are useful for diagnosing and treating an related disorder, e.g. systemic lupus crythematosus, rheumatoid arthritis. The present sequence represents a DNA encoding a PRO
  New nucleic acid encoding PRO polypeptide, useful for diagnosing and treating an immune related disorder, e.g. systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis, thyroiditis, or diabetes mellitus.
  Antilliammatory; Immune disorder; Dermatological; Immunosuppressive; Antirheumatic; Antiarthritic; Osteopathic; Hemostatic; Antianemic; Antithyroid; Antidiabetic; Nephrotropic; CNS-Gen.; Hepatotropic; Virucide; Gastrointestinal-Gen.; Antipsoriatic; Antiasthmatic; Antialiamic, Gen.; Antipsoriatic; Antiasthmatic;
   oligonucleotides, all
  11-AUG-2003; 2003US-0493546P
  11-AUG-2004; 2004WO-US026249
  Sequence 2537 BP; 756
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   Claim 1; SEQ ID NO 1453; 158pp; English.
   WPI; 2005-182330/19.
  Abbas
  24-FEB-2005
   WO2005016962-A2
  Homo
  05-MAY-2005
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  ADY15647 standard;
  (GETH ) GENENTECH
  Antiallergic;
   DNA encoding a PRO polypeptide, SEQ ID NO 1453
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   2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
  3499
   sapiens.
   ,
  46;
   1 Similarity
46; Conser
   invention
   Similarity
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  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 3544
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   1.5%;
ilarity 100.0%;
Conservative
  Conservative
  (first
  ds;
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  DNA;
   Ouyang
  entry)
   1.5%;
  ?
  유
  6530
  503
  which
   ; Score 46; DB
$; Pred. No. 1e-
0; Mismatches
   Σ
  0;
   Score 46;
Pred. No.
  BP.
  Ç
  are used in the exemplification
  Williams MP,
  473
  Mismatches
  ဂ္
  805
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1e-10;
  1.1e-10;
hes 0;
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  2057
  H
  Wood WI,
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  <u>.</u>
   Length 2537;
   T; 0
  ď;
  Length 6530;
  Indels
  Indels
  0
   U; 0 Other;
  Other;
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  ij
  0
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  얁
  a
   Gaps
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     exon
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  exon
   exon
  exon
   Xey
   ACC82887;
   exon
  Homo sapiens.
                    05-NOV-2002; 2002WO-US035479
   intror
   intron
  WO2003040328-A2
   intron
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  intron
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   intror
  intror
   15-MAY-2003.
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```
Human thyroid hormone receptor interactor 6 (TRIP6) gene
  (first entry)
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Human; antisense; thyroid hormone receptor interactor 6; TRIP6; tumour; OPA-interacting protein-1; OIP-1; zyxin-related protein-1; prophylaxis; inflammation; therapy; hyperproliferative disorder; infection; cancer; chromosome 7q22; ZRP-1; ds. fragment.

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  /number= 2
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Matches 46
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24-FEB-2000
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16-MAR-2000
17-MAR-2000
18-APR-2000
19-MAY-2000
07-JUN-2000
07-JUN-2000
07-JUL-2000
07-JUL-2000
07-JUL-2000
11-JUL-2000
   The invention relates to antisense compounds targetted to a nucleic acid encoding thyroid hormone receptor interactor 6 (TRIP6) to inhibit its expression. TRIP6 is also known as OPA-interacting protein-1 (OIP-1) and zyxin-related protein-1 (ZRP-1). TRIP6 DNA is located on chromosome 7q22. Antisense compounds of the invention are useful for modulating the expression of TRIP6 and for treating diseases or conditions associated with the expression of TRIP6 such as hyperproliferative disorders (e.g. cancer). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. The are also useful in antisense therapy the present segment is pathway.
  02-AUG-2001.
  WO200155320-A2
  Homo
  cancer;
   Human; reproductive system related antigen; reproductive system disorder;
   Human reproductive system related antigen DNA SEQ ID NO:
   21-NOV-2001 (first entry)
   AAL06913;
   AAL06913 standard; DNA; 13409
  Sequence
  Example 15; Page 89-93; 111pp; English.
  New antisense oligonucleotides targeted to nucleic acids encoding thyroid hormone receptor interactor 6, useful for diagnosing or treating hyperproliferative disorders, such as cancer.
  08-NOV-2001; 2001US-00008789.
  17-JAN-2001; 2001WO-US001339.
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
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  2003-430662/40.
  46;
   Similarity
  gene therapy; ds.
  CF, Dobie K;
  ISIS PHARM INC.
  7001 BP; 1380 A; 1954 C; 1984 G; 1683 T; 0 U; 0 Other;
  The present sequence is human TRIP6 gene fragment
; 2000US-0179065P.

2000US-0184664P.

2000US-0184635P.

2000US-018635P.

2000US-0199076P.

2000US-0199076P.

2000US-0205515P.

2000US-020467P.

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2000US-021647P.

2000US-0217487P.
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14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
14-AUG-2000;
   2000US-0217496P
2000US-0218799P
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2000US-023544P
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Matches
                          Query Match
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08-NOV-2000
17-NOV-2000
17-NOV
  The present invention provides the protein and coding sequences of number of human reproductive system related antigens. These can be in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encodiprotein of the invention
  Isolated nucleic acid molecule encoding a reproductive system antigen used in preventing, treating or ameliorating a medical condition.
   Rosen
  Sequence
   Disclosure;
  2001-465570/50.
  Ş
Similarity
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  2000US-0244617P.
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2000US-0246476P.
2000US-0246478P.
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  3673
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   9601; 1297pp +
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   2856
  Sequence Listing; English.
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07-JUL-2000
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14-JUL-2000
14-AUG-2000
15-SEP-2000
01-SEP-2000
  Cytostatic; immunosuppressive; nootropic; neuroprotective; antiviral; antiallergic; hepatotropic; antidiabetic; antiinflammatory; antiulcer; vulnerary; antionvulsant; antibacterial; antifungal; antiparasitic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder;
   17-JAN-2001;
  02-AUG-2001
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2000US-024921P.
2000US-024924P.
   밁
   SEXEXEXEX
   δ
  The invention relates to novel genes (ABA07454-ABA08224) and proteins CC (ABB10743-ABB10980) useful for preventing, treating or ameliorating CC medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. CC The nucleic acids, proteins, antibodies and (ant)agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and CC ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, CC breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune CC disorders e.g. Addison's disease, allergies, autoimmune haemolytic CC anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, CC multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) CC cardiovascular disorders such as myocardial ischaemias; (d) wound healing (i) enurological disease e.g. cerebral anoxia and epilepsy; and (f) (i) enurological diseases e.g. cerebral anoxia and epilepsy; and (f) CC infectious diseases such as viral, bacterial, fungal and parasitic CC infectious diseases such as viral, bacterial, fungal and parasitic confectious diseases such as viral, bacterial, fungal and parasitic confectious diseases such as viral, bacterial, fungal and parasitic confectious diseases such as viral, bacterial, fungal and parasitic confectious diseases such as viral, bacterial, fungal and parasitic confectious diseases such as viral, bacterial, fungal and parasitic confectious diseases such as obtained in electronic format directly form wife of the printed specification, but was obtained in electronic format directly forms.
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Matches 46
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06-DEC-2000;
08-DEC-2000;
   Sequence 13409 BP;
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   Disclosure; SEQ ID
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   WPI; 2001-488786/53.
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  21-MAY-2002
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  Local Similarity
   12121
   3071
  Ş
   46;
   sequence #928 encoding novel human connective tissue polypeptide
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   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
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Gaps

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20-OCT-2000
20-OCT

| 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | X S X X G X X G G G G G G G G G G G G G                                                   |
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| 2000US-0189874P. 2000US-0198123P. 2000US-0198074P. 2000US-021964FP. 2000US-021664FP. 2000US-021664PP. 2000US-021664PP. 2000US-0217496P. 2000US-0217496P. 2000US-0217496P. 2000US-0225214P. 2000US-0225214P. 2000US-0225266P. 2000US-0225267P. 2000US-0225447P. 2000US-022547P. 2000US-023144P. 2000US-02314P. 2000US-02314P. 2000US-02314P. 2000US-02314P. 2000US-02314P. 2000US-02314P. 2000US-02314P. 2000US-02314P. 2000US- | A1.  2001WO-US001322. 2000US-0179065P. 2000US-0180628P. 2000US-0184664P. 2000US-0186350P. |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                                                                           |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                                                                           |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                                                                           |
| ***************************************                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | ס ס ס ס ס ס ס ס ס ס ס ס ס ס ס ס ס ס ס                                                     |
| 13-OCT 2000; 20-OCT 2000; 20-OCT 2000; 20-OCT 2000; 20-OCT 2000; 20-OCT 2000; 20-OCT 2000; 01-NOV 2000; 08-NOV 2000; 17-NOV 2000; 17-NO |                                                                                           |
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  Ś
  Antiarteriosclerotic; immunosuppressive; antirheumatic; antiarthritic;

Wantiinflammatory; antiallergic; antiakhmatic; dermatological;

Wantiinflammatory; antiallergic; antibacterial; antiparastic;

Wanephrotopic; virucide; fungicide; antibacterial; antiparastic;

Wantiinflammatory; ds; connective tissues disorder; rheumatoid arthritis;

Wantiinflammatory; ds; connective tissues disorder; rheumatoid arthritis;

Wantiinflammatory; ds; connective tissues disorder; rheumatoid arthritis;

Wantiinflammatory; disease; parkinson's disease; cardiovascular disease;

Watherosclerosis; myocarditis; cardiopulmonary bypass complication;

Wantiinflammatory; cardition; Crohn's disease; nephritis;

Watherosclerosis; inflammatory condition; Crohn's disease; nephritis;

Wastrointestinal disorder; inflammatory bowel disease;

Watherosclerosliferative syndrome;

Wantiinflammatory condition; inflammatory bowel disease;

Wastrointestinal disorder; HIV; AIDS; infection;

Watheroscome delerification; disorder; HIV; AIDS; infection;
  Matches 46;
   Query Match
Best Local
  The present invention relates to the isolation of novel human connective tissue related polypeptides (ARU86435-ARU86923) and the polynucleotide (CDNA and genomic) sequences encoding them. The sequences of the invention are useful in the diagnosis, treatment, prevention and/or prognosis of diseases associated with connective tissue(s), including cancer. The polynucleotide sequences of the invention are also useful in gene therapy. ABK42102-ABK43116 represent genomic sequences encoding the novel human connective tissue related polypeptides. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
   Nucleic acid encoding novel connec used in diagnosing, preventing, tr as cancer or rheumatoid arthritis.
 31-JAN-2000; 2000US-0179065P.
04-FEB-2000; 2000US-0180628P.
   07-MAR-2002; 2002US-00092154.
  connective tissue related polynucleotide; gene;
   cytostatic; neuroprotective; nootropic; antiparkinsonian; cardiovascular;
  Connective tissue related genomic DNA #928.
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2000US-0231244P
2000US-0231244P
2000US-0231244P
2000US-0231443P
2000US-0231244P
2000US-0231443P
2000US-0231449P
   2000US-0184664P.
2000US-0186350P.
2000US-0189874P.
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02-OCT-2000)
02-OCT-2000)
13-OCT-2000)
20-OCT-2000)
20-OCT-2000)
20-OCT-2000)
20-OCT-2000)
20-OCT-2000)
20-OCT-2000)
08-NOV-2000)
09-DC-2000)
17-NOV-2000)
17-NOV
New conn
for treatissue,
   Rosen
connective tissue-related polypeptides and polymucleotides, useful treating, preventing and/or prognosing e.g. disorders of connective sue, (e.g. rheumatoid arthritis), cancers, cancer metastases and/or
  2003-634869/60
DB; ADB59732.
   HUMAN
   Ruben SM,
  2000US-0246477P
2000US-0246524P
2000US-0246525P
2000US-0246525P
2000US-0246528P
2000US-0246610P
2000US-0246611P
2000US-0246611P
2000US-0246611P
2000US-0246611P
2000US-0249210P
2000US-0249210P
2000US-0249211P
2000US-02492189
2000US-02492199
2000US-0250391P
2000US-0250391P
2000US-02511989
2000US-02511856P
  2000US-0237039P.
2000US-0237040P.
2000US-023993P.
2000US-023993P.
2000US-0241285P.
2000US-0241785P.
2000US-0241786P.
2000US-0241786P.
2000US-024186P.
2000US-0241809P.
2000US-0241809P.
2000US-0241809P.
2000US-0241809P.
2000US-024181P.
2000US-024181P.
  2000US-0251868P.
2000US-0251869P.
2000US-0251989P.
  2001US-00764847.
   GENOME
  SCI
   Barash
   SC
```

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ARBSULT 92
ADC210
XX
ADC210
XX
ADC210
XX
ACA DDC210
XX
ACA DDC210
XX
ACA DC210
XX
A
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  S
   The invention describes an isolated nucleic acid molecule (I), which comprises a sequence that is at least 95 % identical to a connective tissue-related polynucleotide encoding connective tissue antigens (CTA). The polypeptide or polynucleotide is useful for preventing, treating, or ameliorating medical conditions in a mammal. The connective tissue polynucleotides and antibodies are particularly useful for treating, preventing and/or prognosing disorders of connective tissues (e.g. rheumatoid arthritis, discoid and systemic lupus erythematosus, scleroderma, or Sjogren's syndrome), cancers, cancer metastases and/or neoplasias (e.g. leukaemia), neurodegenerative disorders (e.g. atherosclerosis, myocarditis or cardiopulmonary bypass (e.g. atherosclerosis, myocarditis or cardiopulmonary bypass (complications), autoimmune diseases (e.g. systemic lupus erythematosus, rheumatoid arthritis, or multiple sclerosis), allergic reactions (e.g.
   Query Match
Best Local S
Matches 46
   27-MAR-2001;
12-SEP-2001;
12-SEP-2001;
   gene therapy; human; secreted protein; haemopoietic disorder; haematological disorder; anaemia; haemophilia; inflammatory disease; cronn's disease; neophastic disease; canculeukaemia; wound healing; epithelial cell proliferation disorder; immune disorder; autoimmune disorder; asthmatic disorder;
  neoplasias.
   18-DEC-2003
   ADC21019
   New human secreted proteins and nucleic acid molecules, useful for preparing a diagnostic or pharmaceutical composition for diagnosing preventing or treating hematopoietic or hematologic disorders, e.g.
  gastrointestinal
  cardiovascular disorder; atherosclerosis; myocarditis;
infectious disease; HIV; AIDS; endocrine disorder; diabetes;
   Human secreted protein-related
  ADC21019;
   26-MAR-2002;
   WO200292787-A2
  Homo sapiens.
   (HUMA-)
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
  2003-129287/12.
   224
  ξ
  46;
   ဓ္
  Similarity
   HUMAN
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
   standard;
  Ruben
   SEQ
  Conservative
   2001US-0278650P.
2001US-00950082.
2001US-00950083.
  2002WO-US009257
   (first entry)
   GENOME
   Ħ
  SM;
   disorder;
   DNA;
  NO 1916; 248pp;
   1.5%;
   SCI INC
   18501
  duodenal ulcer; gastroenteritis;
   Score 46; DB; Pred. No. 9.9
  <u>,</u>
   BP
   DNA sequence #437.
   English.
  DB 9; ; . 9.9e-11
   Length 18501;
   Indels
  3116
  269
   gene;
   Ö
   Gaps
   cancer;
  g
   0
```

Disclosure;

SEQ

ID NO

973; 1512pp; English

amino acid and coding sequences

0£ human

invention comprises the

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RESULT 93
ABT17021/c
ID ABT17021 standard; DNA; 18501
  Query Match
Best Local :
  Matches
   useful for detecting, preventing, diagnosing, prognosticating, treating or ameliorating: haematopoletic or haematological disorders (e.g. anaemia and haemophilia); inflammatory disorders (e.g. inflammatory bowel disease and Crohn's disease); neoplastic disease (e.g. cancer and leukaemia); wound healing and disorders of epithelial cell proliferation; immune disorders (e.g. autoimmune disorders and asthmatic disorders); cardiovascular disorders (e.g. atherosclerosis and myocarditis); infectious disease (e.g. HIVAIDS); endocrine disorders (e.g. disorders); and gastroenteritis). The present DNA sequence was used in the exemplification of the invention.
                  The invention comprises the amino acid and coding sequences of human secreted proteins. The DNA and protein sequences of the invention are useful for the diagnosis and treatment of allargic disorders, asthmatic disorders and immediate hypersensitivity diseases (e.g. hay fever, allergic conjunctivitis and allergic rhinitis). The proteins of the invention are also useful for identifying a binding partner. The nucleic acids of the invention are also useful for observables.
  Use of human secreted proteins and nucleic acids for preparing a diagnostic or pharmaceutical composition for diagnosing or treating allergic or asthmatic disorders, e.g. asthma, hay fever, or allergic
   27-MAR-2001; 2001US-0278650P
12-SEP-2001; 2001US-00950082
12-SEP-2001; 2001US-00950083
  03-OCT-2002
   WO200277188-A2
   binding partner
  Human; gene; ds; protein therapy; immediate hypersensitivity
   Sequence 18501 BP;
   conjunctivitis
   WPI; 2003-175010/17.
   26-MAR-2002; 2002WO-US009239
  radiation
   allergic disorder;
   Human secreted
   03-APR-2003
   secreted proteins. The DNA and protein sequences of the invention
   (HUMA-) HUMAN GENOME SCI INC
  Local Similarity
   ergic disorder; asthmatic disorder; gene therapy; secreted fever; allergic conjunctivitis; allergic rhinitis; ding partner identification; chromosome identification;
  3071
  ξ
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 18233
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
   hybrid mapping; long-range restriction
   Ruben
  Page 810-815; 823pp; English
  Conservative
   (first
  or rhinitis.
  protein-related DNA sequence - SEQ
   SM;
   5301 A; 3948 C; 3746
   entry)
  100.0%;
   1.5%;
ç
   0;
  Score 46;
Pred. No.
   뫄
  Mismatches
  DB 10; ]
9.9e-11;
   G; 5504 T; 0 U; 2 Other;
  0
   Length 18501;
  mapping
  Indels
   ID No
  or allergic
   <u>,,</u>
  disease;
   protein;
   Gaps
   are
   0
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RESULT 94
ABZ68161/G
ID ABZ68
XX ABZ68

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   8
  SXS
   Matches
  Query Match
   treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant) agonists are useful in the diagnosis, treatment and prevention of:

(a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung or urogenital, (b) immune disorders e.g. Addison's disease, allergies, autoimmune haemolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) infectious diseases such as viral,
  New human secreted proteins encoded by genes contained in cDNA clones (e.g. HGCAC19), useful for preventing, treating or diagnosing e.g. AII multiple sclerosis, herpes virus, leukemia, tick-borne encephalitis or West Nile fever.
   The invention relates to novel human genes (ABZ66891-ABZ68209) and the encoded secreted proteins (ABP99470-ABP99872) useful for preventing,
  antifungal; antiparasitic; cardiant; immune disorder; infection; vaccine;
cardiovascular disorder; neurological disease; nephrotropic;
   antiarthritic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiulcer; anticonvulsant;
  Sequence 18501 BP;
  Disclosure;
  WPI; 2003-040583/03
   12-SEP-2001;
12-SEP-2001;
  27-MAR-2001;
   26-MAR-2002; 2002WO-US009188.
  03-OCT-2002
  WO200277186-A2
   Homo sapiens.
  vulnerary;
  virucide;
  Human; secreted protein; nootropic; neuroprotective;
  Human secreted protein encoding genomic DNA SEQ
  26-MAR-2003
   ABZ68161;
  ABZ68161
   (HUMA-) HUMAN
  Local
  18278
  sequence represents a human secreted protein-related DNA sequence
  therapy;
   ξ
   46;
  Similarity
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
   standard;
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
   dermatological;
                           fungal
   antibacterial;
   Page 2321-2325;
   Conservative
  2001US-0278650P
2001US-00950082
2001US-00950083
   (first entry
  GENOME
  gene;
                           and parasitic infections
   5301 A; 3948 C; 3746
  DNA; 18501
  dg
   SCI
   100.0%;
   1.5%;
  antiparkinsonian; antisickling;
   immunosuppressive;
  2423pp; English
   0;
   Score 46;
Pred. No.
   ВP
   Mismatches
   DB 10; 1
. 9.9e-11;
   Ģ
   5504 T; 0 U; 2 Other;
   antiinflammatory;
   ij
  Length 18501;
   Indels
   ĕ
  cytostatic;
   antianaemic;
   <u>.</u>
  anti-HIV;
   Gaps
   e
R
   0;
```

Human

purinergic receptor P2X4 gene sequence

numan;

gene; ds.

fat deposition; leanness; non-insulin dependent diabetes mellitus; NIDDM; purinergic receptor; P2X4; antidiabetic; anorectic; diabetes; obesity;

```
RESULT 96
ADH36221/c
ID ADH36221 standard; D
XX
AC ADH36221;
XX
DT 11-MAR-2004 (first of the control of the con
  ADI81379/c
ID ADI813
XX
AC ADI813
AC ADI813
XX Y
AC ADI813
XX Human
XX Human
XX Human
XX Homo s
XX Homo s
XX I ISIS-
XX O1-JUL
XX O1-JUL
XX O1-JUL
XX O1-JUL
XX I Dobie
XX WPI; 2
XX WPI; 2
XX WPI; 2
XX The in
CC The in
CC associ
CC associ
CC disord
CC associ
CC associ
CC disord
CC muclei
CC disord
CC Tuclei
CC associ
  닭
   S
  S
  밁
  S
   RESULT 95
   Matches
   Matches
  Query Match
Best Local (
  Sequence 18501 BP; 5301 A; 3948 C; 3746 G; 5504 T; 0 U; 2 Other;
  New antisense oligonucleotides for modulating P2X4 expression, diagnosing, preventing or treating conditions associated with neurological disorders, osteoporosis or rheumatoid arthritis.
  The invention comprises antisense oligonucleotides that are targeted to a nucleic acid encoding P2X4. The antisense oligonucleotides are useful for inhibiting the expression of P2X4 in cells or tissues to treat diseases associated with P2X4 expression, such as: neurological disorders, bone disorders (e.g. osteoporosis), or rheumatoid arthritis. The present nucleic acid represents the human P2X4 genomic DNA sequence.
  WPI; 2004-081656/08
  01-JUL-2002; 2002US-00187659
   01-JUL-2002; 2002US-00187659.
   22-APR-2004
   01-JAN-2004.
   Homo sapiens.
   human;
  antisense oligonucleotide; P2X4; P2X4-associated diseases;
  Human P2X4 genomic DNA sequence
  ADI81379;
   ADI81379
   Sequence 25001 BP; 5940 A; 6387 C; 6220 G; 6454 T; 0 U; 0 Other;
   (ISIS-) ISIS
  US2004002152-A1
  neurological disorder; bone disorder; osteoporosis; rheumatoid arthritis;
  Local
   18278
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
  2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
   46;
   46;
   15;
   Similarity
  Similarity
   standard;
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 18233
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2588
   SEQ ID NO 11; 67pp; English.
   Conservative
   PHARM INC
   (first
                                     (first entry)
   DNA;
   DNA; 28616
  1.5%; U.
100.0%; Pr
   entry)
   1.5%;
  25001
  0;
   Score 46; pred. No.
  Score 46;
Pred. No.
   ВP
  ВP
   Mismatches
   Mismatches
  DB 10; I
9.9e-11;
   DB 12; I
9.8e-11;
  0,
   Length 25001;
  Length 18501;
  Indels
   Indels
  3116
   0
  0
   Gaps
  Gaps
   e.g.
   0
  0
```

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Query Match
Best Local S
Matches 46
  This invention relates to a novel method of diagnosing a predisposition to fat deposition, leanness or non-insulin dependent diabetes mellitus (NIDDM) in a subject. The method comprises detecting the presence or absence of a polymorphic variation associated with fat deposition, leanness or NIDDM at a polymorphic site in a purinergic receptor (P2X4) nucleotide sequence in a nucleic acid sample from a subject. The invention may be useful for the development of compounds with an antidiabetic or anorectic activity. The method is useful for diagnosing a predisposition to fat deposition, leanness or NIDDM. The nucleic acid encoding the polymeptide is useful for diagnosing conditions or diseases including fat deposition or NIDDM, also in treating diabetes and obesity. The present sequence is that of the purinergic receptor (P2X4) nucleotide sequence which was used in the method of the invention.
  Diagnosing predisposition to fat deposition, leanness or non-insulin dependent diabetes mellitus (NIDDM) comprises detecting the presence absence of a polymorphic variation in a purinergic receptor.
   variation
  variation
   variation
   variation
   variation
  Homo sapiens
  Adam GIR, Langdown ML,
  04-JUN-2002; 2002US-0386012P
  04-JUN-2003; 2003WO-US017676
  11-DEC-2003
  variation
  Sequence 28616 BP; 6868 A; 7260 C; 7008 G; 7438 T; 0 U;
  Claim 1; SEQ ID NO 1; 154pp; English.
  P-PSDB;
  variation
  (SEQU-) SEQUENOM INC.
  46;
                 Similarity
  ADH36222, ADH36223.
  Conservative
  /standard name= "Single nucleotide replace(22713,T)
   /standard name= "Single nucleotide polymorphism"
replace(14744,T)
   /standard name= "Single nucleotide polymorphism"
replace(21708,G)
   /standard_name= "Single nucleotide polymorphism"
replace(15847,T)
   /standard name= "Single nucleotide polymorphism"
replace(15059,G)
   Location/Qualifiers replace(11030,T)
   /standard name= "Single nucleotide polymorphism"
replace(17338,T)
  standard_name=
   *tag=
  *tag=
                1.5%;
  Roth
  0
  Score 46; DB
Pred. No. 9.7
0; Mismatches
  RB,
  "Single nucleotide polymorphism"
  Denissenko
                 DB 12; 1
. 9.7e-11;
   ጃ
ማ
,
                             Length 28616;
     Indels
  Smylie
  polymorphism"
   42
   Other;
    Gaps
   õ
     0
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2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940

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4783

GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 4738

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RESULT 97
ADZ42284/c
ID ADZ42284 standard;
  exon
   exon
This invention relates to a novel method for testing hypertensive renal
                Claim 1; SEQ ID NO 11; 440pp; Japanese.
                                       Testing hypertensive renal disease factor, by determining polymorphism in genotype of gene relevant to hypertensive renal disease, and estimating risk factor for hypertensive renal disease based on determined genotype,
  WPI; 2005-326228/34.
  09-OCT-2003; 2003JP-00350959
  09-OCT-2003; 2003JP-00350959
  JP2005110606-A
  3'UTR
   variation
   exon
  exon
   CDS
   5'UIR
  Key
  single
  Human Klotho gene with C49620T SNP Seq 11.
   ADZ42284;
  intron
  intron
  intron
  Homo sapiens
   renal disease;
   14-JUL-2005
  disease; nephrotropic; SNI
nucleotide polymorphism;
  KOKURITSU JUNKANKI BYO CENT SOCHO.
DOKURITSU GYOSEI HOJIN IYAKUHIN IRYO KIK.
   (first
  49620
   /standard
52961. .54
  43821
  /*tag=
   49453. .50554
/*tag= i
  44141. .49452
  number=
  /*tag=
  number=
   13052. .43820
   12541. .43051
  /*tag= c
/product= "Klotho
   location/Qualifiers
  number= 4
  number=
  *tag=
   number= 2
   number=
   *tag=
   number= 1
  *tag=
   16. .6034
  DNA;
   entry)
   .54916
k
  .42540
  .50554
   .44140
  е
  ω
  58000
  name=
   SNP detection;
sm; SNP; Klotho;
  ВP
  "Single nucleotide polymorphism"
  protein"
   ds;
  gene
```

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밁
  S
  1.5%;
Best Local Similarity 100.0%;
Matches 46; Conservative
to a method for treating cancers involving administering to a patient an inhibitor of CAP, and a method of screening for anticancer activity in a potential drug involving providing a cell that express a CA gene, contacting a tissue sample derived from a cancer cell with an anticancer drug candidate and monitoring the effect of the anticancer drug candidate on expression of the CA gene. The CAP proteins are useful for detecting cancer associated with expression of a CAP protein in a test cell sample and for screening for a bioactive agent capable of modulating the activity of a CAP protein. The CA nucleic acids are useful for diagnosing
  adrenoreceptor, aldosterone synthetase, endothelium nitrogen monoxide synthetase, klotho and a sodium-calcium exchanger. Furthermore, it provides primers and probes for determining hypertensive renal disease factors, in particular in relation to renal diseases including hypertensive early renal disease and hypertensive kidney blood flow obstruction. The method enables detection of risk factors, and thus helps in preventing or delaying renal disease. This polymucleotide sequence is in preventing or delaying renal disease. This polymucleotide sequence is the control of the the contro
   genotype of a gene relevant to hypertensive renal disease and estimating the risk factor for developing the disease accordingly. The present invention describes identifying gene polymorphisms in at least one of the following genes, namely endothelin converting-enzyme 1, mineralocorticoid receptor, urotensin II, superoxide-dismutase 3, thiazide sensitivity NaCl symporter, guanosine cyclase-A, hepatocyte growth factor, beta-3
   The invention relates to cancer-associated proteins (CAP) and the cancer-associated (CA) nucleic acids encoding them. The invention also relates
  Novel human cancer associated protein encoded within open reading of cancer associated gene, useful as targets for diagnosing cancer
  WPI; 2004-499109/47
  Human cancer-associated (CA) gene HD07-076.
   18-NOV-2004
   ABD33407;
  ABD33407
   Sequence
  Claim 16; SEQ ID NO 526; 182pp; English.
   Morris DW,
  17-DEC-2002; 2002US-00322281
  15-DEC-2003;
   15-JUL-2004
  WO2004058146-A2
  Homo sapiens.
   ds; cancer; cytostatic.
   Human; cancer-associated protein; CAP; cancer-associated gene; CA; gene;
  disease.
   (SAGR-) SAGRES DISCOVERY INC
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  58000 BP; 16432 A; 11653 C; 11941 G; 17973 T; 0 U; 1 Other;
  standard; DNA; 58922 BP.
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
   Specifically, it refers to determining polymorphisms
   Malandro
  2003WO-US040081
   (first entry)
   NS;
  <u>.</u>
  Score 46;
Pred. No.
   Mismatches
  DB 14; I
. 9.5e-11;
   Length 58000;
  Indels
   36151
   cancer.
  0
   in the
  Gaps
   frame
  0
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CC associated (CA) nucleic acid, comprising two or more nucleic acid probes.

CC The invention also relates to a peptide array comprising two or more for the invention also relates to a peptide array comprising two or more compound that binds to a polypeptide, an isolated antibody or its fragment which composition comprising the polypeptide or its antigen binding comprising the polypeptide or its antipen binding composition comprising the host expressing antibodies against the antigen or its antigen binding fragment, a composition comprising the composition or a carrier, a method of screening for anticancer activity, a composition comprising the composition of a carrier, a method of diagnosing cancer, a composition of a carrier, a method of it reating cancer and a method of inhibiting expression of a CA nucleic acids in a cell. The CA nucleic acids are useful for detecting CA composition in a cell. The CA nucleic acids are useful for detecting CA composition of a cancer cells in an individual which involves contacting cells crome the individual with the antibody and detecting a complex of a CA composition is useful for inhibiting growth of cancer cells in an individual or for delivering a therapeutic agent to cancer cells in an individual or for delivering a therapeutic agent to cancer for treating carcer and for inhibiting growth of cancer for treating carcer and for inhibiting a complex of a cancer for treating carcer and for inhibiting and the antibody.
  Matches
   Query Match
Best Local
   cancer, involving determining the expression of a CA nucleic acid in a tissue. This sequence represents a human CA gene of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
   The invention relates to a nucleic acid array for detecting a cancer associated (CA) nucleic acid, comprising two or more nucleic acid pr
   Disclosure; SEQ ID NO 60; 198pp; English
   comprises two or more nucleic acid probes.
  Nucleic acid array useful for detecting cancer associated nucleic
  Diagnosis;
   Sequence 58922 BP; 13257 A; 15256 C; 16255 G; 14082 T; 0 U; 72
   Morris DW,
   23-SEP-2003; 2003US-00669920
   07-APR-2005.
  WO2005031001-A2
   Homo sapiens
  cytostatic;
   Human cancer-associated genomic DNA #7.
  16-JUN-2005
   ADZ12540;
   ADZ12540 standard; DNA; 70271 BP
  (CHIR )
  23-SEP-2004; 2004WO-US031617
   Local Similarity
   ftp.wipo.int/pub/published_pct_sequences
  53826 CAAGATTGTGCCÁCTGCÁCTCCAGCCTGGGCÁACAGÁGCÁAGACTC 53781
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  46;
  CHIRON CORP
   DNA microarray; microarray; biochip; cancer;
   Malandro
  gene;
  Conservative
  (first
  ds.
  entry)
  1.5%;
   MS.
   cancer and
   Score 46; DB; Pred. No. 9.5
   °.
   for inhibiting expression of
   DB 13;
9.5e-11;
   0
  Length 58922;
   Indels
   neoplasm;
   0
   a
Ç
   Other;
   Gaps
gene
   0
```

RESULT 101

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RESULT 100
ADE95974
  ឧឧ
밁
                          8
  片
   S
   Matches
  Matches
  Query Match
   Query Match
Best Local Similarity
   diagnosis and treatment of cancer, especially carcinomas, as well as the use of compositions in screening methods. The compositions of the invention may have cytostatic activity whilst the disclosed sequences may be useful for gene therapy. The carcinoma associated nucleic acids and proteins are useful for diagnosing and treating carcinomas, for example lymphoma, breast cancer, prostate cancer or leukaemia, or for screening drug candidates or bloactive agents capable of binding to, or modulating the activity of, a carcinoma associated protein. The present sequence is the genomic DNA sequence of the human SYK gene which is a carcinoma associated gene of the invention.
  Sequence
   New carcinoma associated nucleic acids and proteins, useful drug candidates, or for diagnosing and treating carcinomas, lymphoma, breast cancer, prostate cancer or leukemia.
   cancer diagnosis; cancer treatment; carcinoma; cytostatic; gene therapy;
lymphoma; breast cancer; prostate cancer; leukaemia; ds; human; SYK.
   Human SYK gene genomic DNA sequence
   12-FEB-2004
   ADE95974;
  ADE95974
  Sequence
  This invention relates to novel recombinant nucleic acids for use
  Claim 1; SEQ ID NO 232; 793pp; English
  WPI; 2003-441462/41.
   Morris DW, Engelhard EK;
   08-NOV-2001; 2001US-00052482
  08-NOV-2002; 2002WO-US036071.
  15-MAY-2003
  Homo sapiens
  WO2003039484-A2
  Local
  cell. This sequence represents human cancer-associated genomic DNA
57089
  42048 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 42093
  invention
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
   46;
  46;
  Similarity
   SAGRES DISCOVERY
  70271 BP; 19379 A; 15870 C; 15381 G; 19641 T; 0 U; 0 Other;
                              CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  96594 BP; 27524 A; 20558 C; 21159 G;
  standard;
 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 57134
   (first entry
   Conservative
  Conservative
  DNA; 96594
  100.0%;
   100.0%;
  1.5%;
  1.5%;
   0
  0
   Score 46;
Pred. No.
  Score 46; DB 10;
Pred. No. 9.3e-11;
   ΒP
   Mismatches
  Mismatches
   DB 14;
9.4e-11
  26914 T;
  Length 96594;
  Length
   Indels
   Indels
  0 U; 439 Other;
   3116
  for screening e.g.
   0
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  Gaps
   Gaps
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  0
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á
 BXS
  밁
   ADB72464
  RESULT 102
   The invention relates to recombinant carcinoma associated (CA) nucleic CC acid sequences from mouse and human (AAA01492-ADA03094), and to CC carcinomia associated proteins (CAP) encoded by them. The CC invention also encompasses expression vectors and host cells comprising a CC CA nucleic acid, a polypeptide (especially an antibody) that specifically companies to the protein, and a biochip comprising CA nucleic acid or CC fragments thereof: The sequences of the invention were identified using CC oncogenic retroviruses, which insert into the genome of the host organism CC at random. Many of these do not carry transduced host oncogens or pathogenic trans-acting viral genes, meaning that cancer incidence is a Cd irect consequence of the effects of proviral integration into host CC protooncogenes. The CA nucleic acid sequences can be used to diagnose CC carcinoma (especially breast cancer, prostate cancer, lymphoma or CC leukaemia) or a propensity to carcinoma by determination of the sequence CC of a CA gene, or by determination of CA gene expression in particular CC tissues. CA nucleic acids, proteins and antibodies are also useful as CC therapeutic agents and in screening and evaluating drug candidates. The CC present sequence represents a specifically claimed human CA nucleic acid sequence of the invention. Note: The complete sequence data for this patent did not form part of the printed specification, but was obtained of in electronic format directly from WIPO at
  Query Match
Best Local Similarity
Matches 46; Conserv
ADB72464;
                               ADB72464 standard; DNA; 96595 BP.
   Sequence
  Claim 1; SEQ ID NO 1244; 245pp; English
   New recombinant nucleic acid encoding carcinoma associated useful for preparing compositions for treating carcinomas.
  WPI; 2003-587068/55
  26-DEC-2002; 2002WO-US041414.
  WO2003057146-A2
   Homo sapiens
  prostate; lymphoma; leukaemia; cytostatic; gene therapy; drug screening;
   Human; carcinoma associated; oncogene; carcinoma; cancer; breast;
  Human SYK carcinoma
   06-NOV-2003
  ADA02726;
  ADA02726 standard;
   ftp.wipo.int/pub/published_pct_sequences.
   Morris
  26-DEC-2001; 2001US-00035832
   17-JUL-2003
  (SAGR-) SAGRES DISCOVERY
   57090
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 57135
   96595 BP;
  Conservative
  (first
   27524 A; 20559 C; 21158 G; 26915 T; 0 U;
  DNA;
   associated gene,
   entry)
   100.0%;
   1.5%;
  96595
  <u>.</u>
   Score 46;
; Pred. No.
  Mismatches
  SEQ ID NO:1244
   DB 9; Le
  Length 96595;
  Indels
   protein,
   439 Other;
  0
  Gaps
  0
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  Ś
   AD056274/c
ID AD056
   RESULT 103
  Matches
  Query Match
Best Local Similarity
  The invention relates to a novel recombinant nucleic acid comprising a nucleotide sequence selected from any of the 660 sequences fully defined in the specification. A polynucleotide of the invention has cytostatic activity, and may have a use in gene therapy, or in a vaccine. The recombinant nucleic acids and polypeptides are useful for treating carcinomas, e.g. lymphomas, cancers, neoplasm, adenocarcinoma, and sarcomas. The present sequence represents a human gene of the invention.
   23-OCT-2001;
08-NOV-2001;
30-NOV-2001;
20-DEC-2001;
   gene therapy; human; ds; gene; melanoma;
melanoma associated polymorphic variation; SNP;
single nucleotide polymorphism; cyclin-dependent
   carcinomas, e.g. lymphomas, cancer, .---- human gene sarcomas. The present sequence represents a human gene
variation
  Homo sapiens
   Human cyclin-dependent kinase 10,
   ADO56274;
   AD056274
  Sequence
   Claim 1; SEQ ID NO 292; 2304pp; English.
   human; ds; cytostatic; gene therapy; vaccine; carcinoma; lymphomas; cancer; neoplasm; adenocarcinoma; sarcoma; gene.
  variation
  12-AUG-2004
  New recombinant nucleic acid, useful for treating carcinomas, lymphomas,
  Morris DW,
  02-MAR-2001;
   26-DEC-2001; 2001WO-US051291.
  WO2003008583-A2
  Homo sapiens
  04-DEC-2003
  (SAGR-) SAGRES DISCOVERY
  57090 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 57135
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGAGCTC 3116
   SYK gene
   46;
   standard; DNA; 99100
  96595 BP; 27524 A; 20559 C; 21158 G; 26915 T; 0 U; 439
   neoplasm,
  1.5%;
ilarity 100.0%;
Conservative
   ; 2001US-00004113.
; 2001US-00052482.
; 2001US-00997722.
; 2001US-00034650.
  Engelhard EK;
  2001US-00798586.
   (first entry)
  (first entry)
/note=
424
  Location/Qualifiers
                             /*tag=
   adenocarcinoma, or sarcomas.
a
"Single nucleotide polymorphism"
   0;
  Score 46;
Pred. No.
   뫈
   Mismatches
  CDK10, genomic sequence
  DB 10; Length 96595; 9.3e-11;
```

0;

Indels

0;

Gaps

0,

Other;

kinase 10; CDK10

| variation 21338 /*tag=                                    | variation | variation 19488 /*tag= w /note= "Single nucleotide variation 20864     | variation /note= "/note= "/not | variation 18077  /*tag= t /note= "Single nucleotide variation 18435 /*tag= " | variation | variation 16586 /*tag= /note=' variation 16824 | /*tag= o /note= "Single nucleotide variation 16275 /*tag= p | /note= " variation                                                                          | variation 9640 /*tag= 1 /note= "Single nucleotide variation 13285 /*tag= m | variation 8081 /*tag= j /note= "Single nucleotide variation 8194 /*tag= k /note= "Single nucleotide                                      | 4647 /*tag= h /note= "Single nucleotide 7960 /*tag= i /note= "Single nucleotide | variation 3525 /*tag= f /note= "Single nucleotide variation 4165 /*tag= g /note= "Single nucleotide | variation 3166 /*tag= d /note= "Single variation 3501 /*tag= e /note= "Single | /*tag= b /note= "Single nucleotide 2898 /*tag= c /note= "Single nucleotide                                                               |
|-----------------------------------------------------------|-----------|------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------|-----------|------------------------------------------------|-------------------------------------------------------------|---------------------------------------------------------------------------------------------|----------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------|
| variation 47754<br>/*tag= ax<br>/note= "Single nucleotide |           | /*tag= au<br>/note= "Single nucleotide<br>variation 44986<br>/*tag= av | variation 44029 /*tag= at /note= "Single nucleotide variation 44692                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            | variation                                                                    |           | variation                                      | variation                                                   | variation 40292  /*tag= al  /note= *Single nucleotide  variation 4069  /*tag= am  /*tag= am | ation                                                                      | FT /*tag= ah FT /note= "Single nucleotide polymorphism" FT variation 34152 FT /note= "Single nucleotide polymorphism" FT variation 39455 | /*tag=<br>/note=<br>variation 27859<br>/*tag=<br>/note=<br>variation 33527      | variation<br>variation                                                                              | /*tag=<br>/note=<br>ation 22081<br>/*tag=<br>/note=<br>ation 23427            | FT /note= "Single nucleotide polymorphism" FT variation 21343 FT /note= aa FT /note= "Single nucleotide polymorphism" FT variation 21599 |

|                                                                                                                                                                                                                                                                            | EXSXXXXXXXX                                                      | PESC ACX                                                                                              | P P P P P P P P P P P P P P P P P P P                                                        |                                                                               |                                                                                                                                                                                                                                                                                   | 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7                                  |
|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------|
| FT variation                                                                                                                                                                    | Human cyclin-d melanoma; DNA single nucleot Homo sapiens. Kev    | 3071 CAAGJ<br>    <br>16521 CAAGJ<br>ILT 104<br>30723/c<br>ADX80723 stand<br>ADX80723;<br>05-MAY-2005 | r variation r variation r variation r variation guery Match Best Local Simila Matches 46; CC | variation variation variation                                                 | variation variation variation variation                                                                                                                                                                                                                                           | variation<br>variation                                                 |
| /*tag= a /*tag= a /*tag= a /*standard_name= "Single nucleotide polymorphism" 519 /*tag= b /*standard_name= "Single nucleotide polymorphism" 2993 /*tag= C /*tandard_name= "Single nucleotide polymorphism" 3261 /*tag= d /*standard_name= "Single nucleotide polymorphism" | pi.<br>Ođej                                                      | CAAGATTGTGCCACTGCACCTGGGCAACAGAGCAAGACTC 3116                                                         |                                                                                              |                                                                               | /*tag= az /note= "Single nucleotide polymorphism" 50476 /*tag= ba /note= "Single nucleotide polymorphism" 50525 /*tag= bb /note= "Single nucleotide polymorphism" 50621 /*tag= bc /note= "Single nucleotide polymorphism" 53410 /*tag= bd /note= "Single nucleotide polymorphism" | 47914<br>/*tag= ay<br>/note= "Single nucleotide polymorphism"<br>49672 |
|                                                                                                                                                                                                                                                                            |                                                                  | <del></del>                                                                                           |                                                                                              |                                                                               |                                                                                                                                                                                                                                                                                   |                                                                        |
|                                                                                                                                                                                                                                                                            |                                                                  |                                                                                                       |                                                                                              |                                                                               | 1                                                                                                                                                                                                                                                                                 | er e                               |
| variat<br>variat<br>variat                                                                                                                                                                                                                                                 | FT variation FT variation FT variation FT variation FT variation | variat<br>variat<br>variat                                                                            | FT variation FT variation FT variation FT variation FT variation FT variation                | FT variation FT variation FT variation FT variation FT variation FT variation | variat<br>variat<br>variat                                                                                                                                                                                                                                                        | FT variation FT FT variation                                           |

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23522
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   *tag= aw
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   standard_name=
  *tag= as
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RESULT 105
AD097818
ID AD0978
XX AD0978
XX AD0978
XX AD0978
XX Cytost
XX Cytost
XX W02004
XX W02004
XX 22-ULL
XX 22-DEC
XX 27-DEC
XX WPI; 2
XX Concert
XX Claim
XX Claim
XX Claim
   밁
  ঠ
   Query Match
Best Local Similarity
Matches 46; Conserv
                                     New isolated cancer associated nucleic acids comprising at least contiguous nucleotides, useful for diagnosing, preventing and/or cancers such as leukemia and lymphoma.
   Homo sapiens
   Human cancer associated sequence HD11-002, SEQ ID 795
   07-OCT-2004
   ADQ97818
  ADQ97818 standard; DNA; 109661 BP
   variation
   variation
   variation
   variation
   variation
   variation
                   Claim 1;
   WPI; 2004-543781/52
  27-DEC-2002; 2002US-00330773
   22-DEC-2003; 2003WO-US041389
   22-JUL-2004
   WO2004060304-A2
  Cytostatic; Gene
   variation
  variation
  variation
   Morris
  (SAGR-)
  16616 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 16571
   3071
   DW, Malandro
  SAGRES DISCOVERY
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
                   SEQ ID
   Conservative 0;
   (first
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/standard_name=
50704
  /*tag= bh
/standard_name=
50608
                    NO 795;
  50260
   /standard_name=
47837
   /standard_name=
   50559
  19992
  standard_name=
   Therapy; cancer; leukemia; lymphoma; Human;
   standard_name=
   *tag= bj
  standard_
  standard_name=
  *tag=
   *tag= bc
standard_name=
  *tag=
   *tag=
   standard_name=
   entry)
   MS.
  1.5%;
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  þf
                  199pp; English
  name=
  Score 46; DB 14;
Pred. No. 9.3e-11;
0; Mismatches 0;
  "Single nucleotide polymorphism"
   "Single
   "Single nucleotide polymorphism"
  "Single
   "Single
   "Single
   "Single
   "Single
  "Single
  "Single
  nucleotide polymorphism'
   nucleotide polymorphism*
  nucleotide polymorphism'
  nucleotide polymorphism"
   nucleotide polymorphism"
  nucleotide polymorphism'
  nucleotide polymorphism"
  nucleotide polymorphism"
   Length 99250;
  Indels
  3116
  0
   g
  Gaps
  : 10
: treating
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present invention relates to cancer associated sequences (ADQ97025-

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RESULT 108
ADZ13631_0
WP Sequence
WP Frag
WP ADZ1
WP ADZ1
WP ADZ1
WP ADZ1
WP ADZ13(
XX AC ADZ13(
XX DT 16-JUN
DE Human
   RESULT 106
ADG70447_1/c
Continuation
   8
  និងនិងនិង
និង
   ABZ79565_1/c
Continuation (2
   밁
  8
   밁
   片
   S
  RESULT
  Query Match
Best Local S
Matches 46
  P Sequence split into P Fragment Name P Fragment Name IP ABZ79565 0 P ABZ79565 2 P ABZ79565 3 P ABZ79565 4
   Query Match
Best Local S
Matches 46
  Matches
   Query Match
  ntinuation (2 of 5) of
Sequence split into 5
  Sequence split into
  Fragment Name
ADZ13631 0
ADZ13631 1
ADZ13631 2
ADZ13631 3
ADZ13631 4
ADZ13631 standard; 1
   ADQ98004). The sequences are useful for the diagnosis, prevention and/or treatment of cancer, such as leukemia and lymphoma. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formate directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
   Sequence 109661 BP; 30680 A; 19371 C; 19986 G; 35350 T; 0 U; 4274 Other;
                       16-JUN-2005
  ADZ13631;
   Pragment Name
ADG70447 0
ADG70447 1
ADG70447 2
ADG70447 3
ADG70447 3
ADG70447 4
  Local Similarity
   Local Similarity
   Local Similarity
   40304
   97599
   97599
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGAGCTC 3116
   2895
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
   46;
   46;
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 97554
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 40349
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
   of 5)
   Conservative 0;
   Conservative
  Conservative
   into
                       (first
  of.
   ហ
   v
  fragments
Begin
   1.5%;
  ABZ79565 f
fragments
Begin
   ADG70447 from base 100001 (Human ANGE-CLLD8-CLLD7 hybrid gene. fragments LOCUS ADG70447 Accession Adg70447
                       entry)
   1.5%;
  100001
200001
300001
   100001
200001
300001
400001
  100001
200001
300001
400001
  400001
   Begin
  1.5%;
   420555
   0;
   Score 46; DB; Pred. No. 9.3
  <u>,</u>
  Score 46;
Pred. No.
   Score 46;
Pred. No.
  from base
   End
110000
210000
310000
410000
   End
110000
210000
310000
  110000
210000
310000
410000
  410000
410846
  ΒP
  LOCUS ADZ13631
  410846
  POCUS
   420555
   Mismatches
   Mismatches
  se 100001 (CLLD8 and NY-REN-34 encoding DNA. ABZ79565 Accession Abz79565
   DB 12; I
. 9.3e-11;
  DB 10; 1
9.3e-11;
  DB 10;
9.3e-11;
   0;
   Accession Adz13631
  0
   Length 110000;
  Length 110000;
   Length 109661;
   Indels
  Indels
   Indels
  97554
  0,
   0
  0,
  Gaps
  Gaps
  Gaps
  0
   0
  0,
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CC The invention also relates to a peptide array comprising two or more consolated polypeptides encoded by a CA nucleic acid sequence, a compound countries to a polypeptide, an isolated antibody or its fragment which countries are a compound countries to a polypeptide, which is prepared by immunizing a host animal collecting cells from the host expressing antipen binding countries arrived or its antigen binding fragment, a composition comprising the polypeptide or its antipen binding fragment, a composition comprising the countries against contribute and a carrier, a method of screening for anticancer activity, a comethod of detecting a CA nucleic acid, a method of diagnosing cancer, a comethod of tracting cancer and a method of inhibiting expression of a CA concleic acids. The antibody is useful for detecting the presence or cabsence of cancer cells in an individual which involves contacting cells from the individual with the antibody and detecting a complex of a CA composed from the cancer cells in an individual which involves contacting cells in an individual. The composition is useful for inhibiting growth of cancer cells in an individual or for delivering a therapeutic agent to cancer cells in an individual. The invention is also useful for diagnosing cancer, for treating cancer and for inhibiting expression of a CA gene in a call. This sequence represents human cancer-associated genomic DNA of the calls in the cancer cepresents human cancer-associated genomic DNA of
   Diagnosis; DNA microarray; microarray; biochip; cancer; neoplasm; cytostatic; gene; ds.
  Disclosure; SEQ ID NO 1151; 198pp; English.
  comprises
   Nucleic acid array useful for detecting cancer associated nucleic acid, comprises two or more nucleic acid probes.
  Morris DW,
   23-SEP-2003; 2003US-00669920
   07-APR-2005
  Homo sapiens.
  associated
  23-SEP-2004;
   WO2005031001-A2
   (CHIR ) CHIRON CORP
invention.
   (CA) nucleic acid, comprising two or more nucleic acid
  Malandro
  2004WO-US031617
  relates to a
  SW
  nucleic acid array for detecting a cancer
   probes
```

Sequence 420555 BP; 131028A; 77271C; 78657G; 131031T; OU; 25680ther;

1.5%;

DB 14; 9.3e-11;

Length 110000;

0

밁 δ Query Match Best Local S Matches 46 Local Similarity nes 46; Conserv 50592 3077 TGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122 Conservative ( k; Score 46; DB k; Pred. No. 9.3 0; Mismatches 0; Indels 0 Gaps

REFERENCE SAN RESULT 109
ADZ13620\_0
WP Sequence split into ADZ13620; ADZ13620 Fragment Name
ADZ13620 0
ADZ13620 1
ADZ13620 2
ADZ13620 3
ADZ13620 3
ADZ13620 4 standard; ຜ DNA; 420555 fragments 100001 200001 300001 400001 Begin 110000 210000 310000 410000 420555 BP. End LOCUS ADZ13620 Accession Adz13620

Human cancer-associated genomic DNA #99

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RESULT 110
ADZ13747 2/c
ADZ13747 2/c
Continuation (3 of 4) of 40 o
   吊
   S
  The invention relates to a nucleic acid array for detecting a cancer CC associated (CA) nucleic acid, comprising two or more nucleic acid probes. CC The invention also relates to a peptide array comprising two or more CC isolated polypeptides encoded by a CA nucleic acid sequence, a compound CC binds to a polypeptide, which is prepared by immunizing a host animal CC with a composition comprising the polypeptide or its fragment which CC that antigen or its antigen binding fragment, a composition comprising the polypeptide or its antipen binding CC fragment and collecting cells from the host expressing antibodies against the antipen or its antigen binding fragment, a composition comprising the CC antibody and a carrier, a method of screening for anticancer activity, a method of detecting a CA nucleic acid, a method of diagnosing cancer, a CC method of treating cancer and a method of inhibiting expression of a CA CC nucleic acids. The antibody is useful for detecting the presence or CC absence of cancer cells in an individual which involves contacting cells from the individual with the antibody, and detecting a complex of a CA CC protein from the cancer cells and the antibody, where the detection of the complex correlates with the presence of cancer cells in the composition is useful for inhibiting growth of cancer cells in an individual. The invention is also useful for dagnosing cancer for treating cancer and for inhibiting expression of a CA gene in a cell. This sequence represents human cancer-associated genomic DNA of the invention is also useful for dagnosing cancer.
P Sequence aplit into 4

P Fragment Name
P ADZ13747 0

P ADZ13747 1

P ADZ13747_2
  Matches
   Query Match
Best Local
   07-APR-2005.
  Diagnosis; DNA microarray; microarray; biochip; cancer; neoplasm;
  16-JUN-2005
  Disclosure;
   WPI; 2005-273395/28.
  Morris
   23-SEP-2003; 2003US-00669920
   23-SEP-2004; 2004WO-US031617
   WO2005031001-A2
   Homo sapiens.
  cytostatic; gene;
   Sequence 420555 BP; 131028A; 77271C; 78657G; 131031T; 0U; 25680ther;
   comprises two or more nucleic
   Nucleic acid array useful for detecting cancer associated nucleic
  (CHIR ) CHIRON CORP
  50592
   cancer-associated
  DW.
  Similarity
  TGTGCCACTGCACTCCAGCCTGGGCAACAGAGCCAAGACTCTGTCTC 3122
  TGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 50637
  Malandro
   SEQ ID NO 1140; 198pp; English.
   Conservative
   (first entry)
  of ADZ13747 from base 200001
o 4 fragments LOCUS ADZ13747
         100001
   100.0%;
  Begin
   1.5%; Score 46;
100.0%; Pred. No.
   genomic DNA #98
   0;
   acid
       110000
210000
310000
   Mismatches
   probes.
   DB 14; Length 110000;
   9.3e-11;
  (Human cancer-associated Accession Adz13747
   0,
   Gaps
   acid,
  genomic
   0
  DNA
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RESULT 111
AAD54480
   Æ
  Best Loc
Matches
  Matches
   Query Match
  Query Match
  The invention relates to a method of Identifying p53 pathway modulating agent. The method involves contacting a test agent with an assay system comprising a purified cation C1- cotransporter interactor protein (CIP) polypeptide or polynucleotide, or their functionally active fragment or derivative. The method is useful for identifying modulators of the p53 pathway particularly for identifying agents for treating disorders (e.g. breast cancer) associated with defective p53 function. Modulators of the invention are useful as targets for novel therapeutics. CIP sequences are useful as modifiers of the p53 pathway, and as therapeutic targets for apoptotic or cell proliferation disorders. The invention is useful in
   05-JUN-2001; 2001US-0296076P.
10-OCT-2001; 2001US-0328605P.
15-FEB-2002; 2002US-0357253P.
  Sequence 117962 BP; 27840 A; 32096 C; 30624 G;
   gene therapy. The present sequence is human CIP DNA
  Identifying p53 pathway modulators for treating or diagnosing d with defective p53 function e.g. breast cancer, by providing an system having a purified cotransporter interactor protein (CIP) polypeptide or nucleic acid.
  03-JUN-2002; 2002WO-US017473.
   Homo sapiens.
  Human; p53 pathway; chloride cotransporter interactor protein; CIP; angiogenic disorder; cell proliferation disorder; apoptotic disorder;
  Human CIP DNA
   26-JUN-2003
  AAD54480
  Disclosure; Page 38-101; 123pp; English
  WPI; 2003-175140/17
  Friedman
  12-DEC-2002.
  WO200299055-A2
  (EXEL-) EXELIXIS INC.
  Local
   ADZ13747_3
  госат
   73741 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 73696
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
                                       3071
  46;
  cancer; gene therapy; ds.
  Similarity
  Similarity
  ۲
CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 18443
                                     CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC
  standard;
  Conservative 0;
  Conservative
  Plowman
   (first
  #1.
  DNA; 117962
   1.5%;
   entry)
  9
   300001
  1.5%;
  Belvin
  0,
  Score 46; DB 8; L
Pred. No. 9.3e-11;
  Score 46;
Pred. No.
   365720
  Mismatches
   Mismatches
  Σ,
  Francis-Lang
  DB 14; 1
9.3e-11;
  Length 117962;
  27402 T; 0 U; 0
   Length 110000
  Ä
  Indels
   Indels
  E
   3116
  á
  0
   0
   an assay
  Punke
   Gaps
  Gaps
  RP;
   0
  0
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RESULT 113
ADC87620/c
ID ADC87620;
XX ADC87620;
AC ADC87620;
XX DT 01-JAN-2004 (first of the control of the con
   RESULT 112
ACN43862
  S
  밁
  The present invention relates to novel DNA and protein sequences which Care associated with carcinomas. The sequences are useful for: (1) for greening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of a bioactive agent capable of modulating the activity of CAP; (iv) for carcinoma; (vi) for inhibiting the activity of CAP; (iv) for carcinoma; (vii) for neutralizing the effect of CAP; (vii) for treating carcinoma; (viii) for neutralizing the effect of CAP; (vii) as a biochip; (cx) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining carcinoma or a propensity to carcinoma; and (xi) for capable of capable or useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent CAP which no sequence data was published
   Query Match
Best Local (
  Matches
   ds; human; GPCR; guanosine triphosphate-binding protein coupled receptor;
   Sequence 141463 BP; 40336 A; 28306 C; 29237 G; 43584 T; 0 U; 0
  Claim 1;
  Recombinant nucleic acid useful for diagnosis and treatment comprises a nucleotide sequence.
   WPI; 2003-328604/31
   01-MAR-2002; 2002US-00087192
   28-FEB-2003; 2003WO-US006235
   WO2003073826-A2
   Homo sapiens
   Cytostatic;
  Human genomic sequence hCG21073.
   ACN43862 standard; DNA; 141463 BP
  12-SEP-2003
  18-NOV-2004
  ACN43862;
   (SAGR-) SAGRES DISCOVERY
  73703
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  46;
   Similarity
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 73748
  SEQ ID NO 22; Opp; English.
   1.5%;
ilarity 100.0%;
Conservative (
   carcinoma; lymphoma; cancer; human; gene;
  (first
  (first entry)
  polynucleotide SEQ ID NO:2073.
   DNA; 144792
   entry)
   0;
  Score 46; DB 11;
Pred. No. 9.2e-11;
0; Mismatches 0;
   ₽₽
  0
  Length 141463;
  Indels
  œ,
  0
  Other;
  carcinoma
   Gaps
  0
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RESULT 114
ADL13904/c
ID ADL13904 s
XX
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  8
   Query Match
Best Local (
  Matches
   ds; gene; osteopathic; antiinflammatory; antiarthritic; joint space narrowing; osteophyte development; joint pai osteoarthritis; SNP; single nucleotide polymorphism.
  The invention relates to a novel polymucleotide encoding a guanosine triphosphate-binding protein coupled receptor. (GPCR). A polymucleotide of the invention may have a use in gene therapy. The polymucleotide and polypeptide are useful for preparing a composition for treating a patient in need of increased or suppressed activity or expression of the guanosine triphosphate-binding protein coupled receptor. The protein sequences shown in ADC87618-ADC87623 represent polymucleotide sequences
  New polynucleotide, useful for preparing a composition for treating patient in need of increased or suppressed activity or expression of guanosine triphosphate-binding protein coupled receptor.
            WPI; 2003-559141/52
  Osteoarthritis-associated polymorphic nucleotide #436
   06-MAY-2004
   ADL13904;
   Sequence 144792 BP; 39827 A; 32142 C; 33413 G;
  Suwa M,
                                   Jones KA,
   20-DEC-2001; 2001US-0342603P
   19-DEC-2002; 2002WO-US041225
  WO2003054166-A2
  Homo sapiens.
  Disclosure; SEQ ID NO 2073; 28pp; English.
  WPI; 2003-315783/31.
   (NAAD-)
  18-JUN-2001; 2001JP-00246789
   18-JUN-2002; 2002EP-00013517
   EP1270724-A2
  Homo sapiens
   Match 1.5%; Score 46; Local Similarity 100.0%; Pred. No.
  60793
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  46;
   ç
  NAT INST ADVANCED IND SCI & TECHNOLOGY.
CENT ADVANCED SCI & TECHNOLOGY INCUBATI
   INCYTE GENOMICS
  Аваі К,
  standard; DNA; 164772
                                    Schafer A;
   the invention.
  Conservative
   (first entry)
  Akiyama Y,
  0
  Aburatani H;
   Mismatches
  ے۔۔۔، gene therapy;
پسطال; joint pain;
polymorphism.
  DB 10;
   INCUBATIO
   9.2e-11;
   39310 T; 0 U;
  Length 144792;
  Indels
  60748
   0;
   100
   Gaps
  Other;
  ng a
of the
   0
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Determining susceptibility of an individual to joint space narrowing, osteophyte development and/or joint pain comprises identifying whether the individual has at least one polymorphism in a polynucleotide encoding Disclosure; SEQ ID ŏ 436; 297pp; English

The invention relates to a method of determining susceptibility of an CC individual to joint space narrowing and/or osteophyte development and/or joint pain comprising identifying whether the individual has at least one CC polymorphism in a polymucleotide encoding at least one of the protein CC listed in the specification. The methods, composition and agent are CC useful for modulating the susceptibility of an individual to joint space CC narrowing and/or osteophyte development and/or joint pain that is CC associated with a disease, preferably osteoarthritis. The cell line and CC the non-human animal are useful for screening for an agent for diagnosing CC an individual having susceptibility to joint space narrowing and/or CC osteophyte development and/or joint pain. This sequence corresponds to the polymucleotide encoding a protein listed in the specification. (Note: The sequence data for this patent did not form part of the printed CC specification but was obtained in electronic format directly from WIPO at CC ftp.wipo.int/pub/published\_pct\_sequences).

Sequence 164772 BP; 50645 A; 32137 C; 31960 G; 50022 T; 0 U; 8 Other,

Matches Query Match Best Local : h 1.5%; Similarity 100.0%; Conservative 0; Mismatches Score 46; Pred. No. DB 10; 9.2e-11; . . Length 164772; Indels Gaps 0

유 δ 111876 2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 111831 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940

RESULT 115 ACN44262

ACN44262 standard; DNA; 168821 ₽P

ACN44262

18-NOV-2004 (first entry)

Human genomic sequence hCG18035.

Cytostatic; carcinoma; lymphoma; cancer; human; gene; 88.

Homo

WO2003073826-A2

12-SEP-2003

28-FEB-2003; 2003WO-US006235.

01-MAR-2002; 2002US-00087192.

(SAGR-) SAGRES DISCOVERY

Morris

2003-328604/31

Recombinant nucleic acid useful for diagnosis and comprises a nucleotide sequence. treatment 얁 carcinoma

OBS Ħ NO 622; Opp; English.

The present invention relates to novel DNA and protein sequences which are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of a bioactive agent capable of modulating the activity of CAP; (Iv) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing

RESULT 116
ADL13935
ADL13935
AC ADL139
XX ADL139
XX OSTEODA
XX OSTEO
XX OSTEODA
XX OSTEODA
XX OSTEODA
XX OSTEODA
XX OSTEODA
XX OSTEO
XX OSTEODA
XX OSTEODA
XX OSTEODA
XX OSTEODA
XX OSTEODA
XX OSTEO
XX OSTEODA
X អន្តិ អ 밁 8 Query Match carcinoma; (vi) for inhibiting the activity of CAP; (vii) for treating carcinoma; (vii) for neutralizing the effect of CAP; (ix) as a blochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent US2002182586A1, for which no sequence data was published Determining susceptibility of an individual to joint space narrowing, osteophyte development and/or joint pain comprises identifying whether the individual has at least one polymorphism in a polymucleotide encoding ds; gene; osteopathic; antiinflammatory; antiarthritic; joint space narrowing; osteophyte development; joint pai Sequence 168821 BP; 39588 A; 43389 C; 45655 G; 40189 T; 0 U; 0 03-JUL-2003 Homo sapiens osteoarthritis; Osteoarthritis-associated polymorphic 06-MAY-2004 ADL13935; ADL13935 standard; DNA; 177866 WPI; 2003-559141/52 Jones KA, 20-DEC-2001; 2001US-0342603P 19-DEC-2002; 2002WO-US041225 WO2003054166-A2 protein. 161763 Local 2895 46; Similarity INCYTE GENOMICS GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 161808 Schafer A; Conservative (first entry) SNP; single nucleotide polymorphism. 100.0%; 1.5% 0, Score 46; Pred. No. ВP Mismatches development; joint pain; nucleotide DB 11; 9.1e-11 .1e-11; Length 168821; Indels gene 0, Other; Gaps 0

Disclosure; SEQ ID NO 467; 297pp; English.

The invention relates to a method of determining susceptibility of an CC individual to joint space narrowing and/or osteophyte development and/or CC joint pain comprising identifying whether the individual has at least one CC polymorphism in a polymuclectide encoding at least one of the protein CC listed in the specification. The methods, composition and agent are CC useful for modulating the susceptibility of an individual to joint space CC narrowing and/or osteophyte development and/or joint pain that is CC associated with a disease, preferably osteoarthritis. The cell line and CC in individual having susceptibility to joint space narrowing and/or control to joint space narrowing and/or control to joint space narrowing and/or control to joint pain. This sequence corresponds to CC the polymuclectide encoding a protein listed in the specification. (Note: CC the sequence data for this patent did not form part of the printed CC specification but was obtained in electronic format directly from WIPO at CC ftp.wipo.int/pub/published\_pct\_sequences).

S

Sequence 177866 BP;

```
RESULT 118
ABQ75562
ID ABQ755
XX
AC ABQ755
XX
AC ABQ755
XX
TI 11-NOV
  문
   8
   밁
  ঠ
  RESULT 117
   Query Match
Best Local
   The present invention describes the human SLC5A8 protein (I), which is a cell surface protein. (I) has cyrostatic activity, and can be used in gene therapy. (I) can be used in detecting and treating SLC5A8-associated cancer, e.g. colon cancer, breast cancer, thyroid cancer or stomach cancer. (I) is also useful in screening assays, predictive medicine and in diagnostic and prognostic assays. The human SLC5A8 gene is located on chromosome 12. The present sequence is used in the exemplification of the
   11-NOV-2002
                              ABQ75562;
  Sequence 181257 BP; 53237 A; 35656 C; 35971 G; 56393 T; 0 U; 0 Other;
  Claim 6; SEQ ID NO 2; 207pp; English.
   New SLC5A8 polypeptide, useful for detecting and treating SLC5A8-associated cancer, e.g. colon, breast, thyroid or stomach cancer.
   Markowitz
   05-JUN-2002; 2002US-0386653P
  05-JUN-2003; 2003WO-US018239
   human; SLC5A8; cell surface protein; cytostatic; gene therapy; SLC5A8-associated cancer; colon cancer; breast cancer; thyroid cancer; stomach cancer; cancer; chromosome 12; gene; ds.
   Human SLC5A8 gene SEQ ID NO:2.
   ADF69677
   ADF69677 standard;
   ABQ75562 standard;
  present invention.
  18-DEC-2003
   WO2003104427-A2
   11-MAR-2004
  (UYCA-) UNIV CASE WESTERN RESERVE.
   Local Similarity
   166193
  Local
  99600
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  2004-062348/06
   46;
   Similarity
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 166238
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 99645
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
  Conservative
   Conservative
   (first entry)
   (first entry)
   DNA;
   DNA; 181257
  53227 A;
   100.0%;
  1.5%; Score 46; DB 10; 1 100.0%; Pred. No. 9.1e-11;
  1.5%;
   188888
  ç,
  0
   Score 46;
Pred. No.
  36632 C; 36825 G; 51154 T; 0 U; 28
   ВP
   ₽P
   Mismatches
  Mismatches
   DB 12; !
9.1e-11;
  0
   Length 181257;
   Length 177866;
  Indels
  Indels
  ٥,
  0,
  Gaps
  Gaps
  Other;
  0
  ٥,
```

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Cloning; characterisation; human; cytochrome P450; CYP 27C1; cytostatic; thyromimetic; antidiabetic; antipsoriatic; tuberculostatic; osteopathic; dermatological; antilipaemic; gene therapy; vaccine; Vitamin D; diabetes; vitamin D metabolite deficiency; hyperparathyroidism; hypoparathyroidism; medullary carcinoma; psoriasis; sarcoidosis; tuberculosis; osteomalacia; chronic renal disease; vitamin D dependent rickets; anticonvulsant; fibrogenesis imperfecta ossium; osteofittis fibrosa cystica; osteoporosis; osteopaenia; osteosclerosis; renal osteodystrophy; rickets; steatorrhoea; glucocorticoid antagonism; idiopathic hypercalcaemia; tropical sprue; malabsorption syndrome; cholesterol steroid; lipid metabolic disorder; cappe; ds.
   Human
  related CYP 27C1 clone RP11-30F3 SEQ ID
   NO:21.
```

WPI; 2002-657595/70 Wisniewski 09-FEB-2001; 2001US-0267410P 22-AUG-2002. WO200264765-A2 Homo sapiens. 11-FEB-2002; 2002WO-CA000163.

New nucleic acid molecules encoding cytochrome P450 proteins, human CYP 27C1 and a hybrid homologs from Xenopus laevis, useful for treating diseases related to vitamin D or vitamin D metabolite deficiency, e.g. parathyroidism and diabetes.

Example 1; Fig 1A; 209pp; English.

The present invention describes an isolated nucleic acid molecule (I) CC encoding human cytochrome P450, CYP 27C1, and a hybrid homologue from CC Xenopus laevis. (I) has thyromimetic, antidiabetic, cytostatic, cc antipsoriatic, tuberculostatic, osteopathic, dermatological and cantilipaemic activities, and can be used in gene therapy and in vaccines. CC The nucleic acid molecules, proteins and methods from the present cc invention are useful for treating diseases related to vitamin D or vitamin D metabolite deficiency, e.g. hyper- and hypo-parathyroidism, CC osudohypo-parathyroidism, Secondary hyperparathyroidsm, diabetes, cc disease, hypophosphatemic VDRR, vitamin D dependent rickets, anticonvulsant treatment, fibrogenesis imperfecta ossium, osteititis cf ibrosa cystica, osteomalacia, osteoporosis, osteopaenia, osteosclerosis, renal osteodystrophy, rickets, glucocorticoid antagonism, idiopathic cor cholesterol, steroid and other lipid metabolic disorders. The present cc which is given in an example from the present invention

밁 Ś Query Match
Best Local Similarity
Matches 46; Conserva 142425 2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940 Conservative 1.5%; Score 46; 100.0%; Pred. No. <u>..</u> Mismatches DB 6; 9.1e-11; Length 188888; Gaps

Sequence 188888

BP; 51055 A; 42661 C;

43560 G;

47708 T; 0 U;

3904 Other;

0

RESULT 119 ADL13570/c ID ADL135 XX AC ADL135 ADL13570; ADL13570 standard; DNA; 193672 BP

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밁
  8
   Query Match
Best Local S
Matches 46
   The invention relates to a method of determining susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain comprising identifying whether the individual has at least one polymorphism in a polymucleotide encoding at least one of the protein listed in the specification. The methods, composition and agent are useful for modulating the susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain that is associated with a disease, preferably osteoarthritis. The cell line and the non-human animal are useful for screening for an agent for diagnosing an individual having susceptibility to joint space narrowing and/or osteophyte development and/or joint pain. This sequence corresponds to the polynucleotide encoding a protein listed in the specification. (Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from WIPO at
  Determining susceptibility of an individual to joint space narrowing, osteophyte development and/or joint pain comprises identifying whether the individual has at least one polymorphism in a polynucleotide encoding
  03-JUL-2003.
   ds; gene; osteopathic; antiinflammatory; antiarthritic; gene therapy; joint space narrowing; osteophyte development; joint pain;
  06-MAY-2004
   19-DEC-2002; 2002WO-US041225
  WO2003054166-A2
   Homo sapiens.
   Osteoarthritis-associated
               Homo sapiens
   Cytostatic; carcinoma; lymphoma; cancer; human; gene;
  Human genomic
   18-NOV-2004
  ACN44650
  ACN44650 standard; DNA; 256157 BP
   Sequence 193672 BP;
  ftp.wipo.int/pub/published_pct_sequences).
  Disclosure; SEQ ID NO 102;
   Jones KA,
   (INCY-) INCYTE GENOMICS INC
  20-DEC-2001; 2001US-0342603P
  Local Similarity
  3015
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 3116
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 2970
   Schafer A;
   .larity 100.0%; F
Conservative 0;
   (first entry
   (first entry)
  sequence hCG38672.
   SNP; single nucleotide polymorphism
   43026 A; 54282 C; 51944 G;
  1.5%;
   polymorphic nucleotide #102.
   297pp; English.
  Score 46; Pred. No.
   Mismatches
  DB 10; I
9.1e-11;
  ..
   43718 T; 0 U;
  Length 193672;
  Indels
  0
   702
  Gaps
  0
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  S
   Query Match
Best Local Similarity
Matches 46; Conserv
  The present invention relates to novel DNA and protein sequences which are associated with carcinomas. The sequences are useful for: (1) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of bioactive agent capable of modulating the activity of CAP; (iv) for a bioactive agent capable of modulating the activity of CAP; (iv) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing carcinoma; (vii) for inhibiting the activity of CAP; (vii) for treating carcinoma; (viii) for neutralizing the effect of CAP; (vix) as a biochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent reconnected.
  Sequence 256157 BP; 70370 A; 54568 C; 55511 G; 73304 T; 0 U; 2404 Other;
   Recombinant nucleic acid useful for diagnosis and treatment of carcinoma comprises a nucleotide sequence.
  01-MAR-2002; 2002US-00087192
   28-FEB-2003; 2003WO-US006235
   WO2003073826-A2
   Human cancer-associated (CA) gene HD07-114.
   18-NOV-2004
  ABD33570 standard; DNA; 256157
  US2002182586A1,
   WPI; 2003-328604/31
                                 Morris DW,
   17-DEC-2002; 2002US-00322281
  15-DEC-2003; 2003WO-US040081
  WO2004058146-A2
   Homo sapiens.
   Human; cancer-associated protein; CAP; cancer-associated gene; CA; gene;
   (SAGR-) SAGRES DISCOVERY INC
  15-JUL-2004.
  (SAGR-) SAGRES DISCOVERY
  192749
   cancer;
   2895
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 192794
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
  SEQ ID
                                 Malandro MS;
   cytostatic.
   Conservative
   (first entry)
  NO 1204; Opp; English.
  for which no sequence data was published
  1.5%;
   0
  Score 46;
Pred. No.
   Mismatches
   DB 11;
9e-11;
  0,
  Length 256157;
  Indels
   0,
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WPI; 2004-499109/47

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  S
  RESULT 122
   Matches
  inhibitor of CAP, and a method of screening for anticancer activity in a potential drug involving providing a cell that expresses a CA gene, contacting a tissue sample derived from a cancer cell with an anticancer drug candidate and monitoring the effect of the anticancer drug candidate and monitoring the effect of the anticancer drug candidate on expression of the CAP proteins are useful for detecting cancer associated with expression of a CAP protein in a test cell sample and for screening for a bloactive agent capable of modulating the activity of a CAP protein. The CA nucleic acids are useful for diagnosing cancer, involving determining the expression of a CA nucleic acid in a tissue. This sequence represents a human CA gene of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bloactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of a bloactive agent capable of modulating the activity of CAP; (iv) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing
   Claim
   Recombinant nucleic acid useful for diagnosis and treatment
   01-MAR-2002; 2002US-00087192
   28-FEB-2003; 2003WO-US006235
   WO2003073826-A2
   Homo
   Cytostatic; carcinoma;
   Human genomic sequence hCG17121.
   18-NOV-2004
   ACN44350;
   ACN44350
  Sequence
   Novel human cancer associated protein encoded within open reading frame of cancer associated gene, useful as targets for diagnosing cancer.
   (SAGR-) SAGRES DISCOVERY
   12-SEP-2003.
   192749
  ftp.wipo.int/pub/published_pct_sequences
   2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 2940
  invention relates to cancer-associated proteins (CAP) and the cancer
   present invention relates to novel DNA and protein sequences which
  ciated (CA) nucleic acids encoding them. The invention also relates method for treating cancers involving administering to a patient a
   2003-328604/31
   sapiens.
   1:
   46;
  Similarity
   OBS
  standard; DNA;
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACAT 192794
  256157 BP; 70370 A; 54568 C;
   a nucleotide sequence.
  Conservative
  ID NO 754; Opp; English.
   (first
   IJ
   ĕ
   entry)
  776; 182pp; English.
  gene,
  1.5%;
  lymphoma; cancer; human; gene;
   276276
  <u>,</u>
  Score 46;
Pred. No.
   Mismatches
  55511 G;
  DB 13;
9e-11;
  13;
  for diagnosing
  0
  73304 T; 0
  Length 256157;
  Indels
  U; 2404 Other;
   of,
  0
  Gaps
  0
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   នន្តន្តន្តន្តន្តន្ត
The present invention describes human cancer-associated (CA) nucleotide compension (I). Also described: (1) an expression vector comprising (I); (C) a host cell comprising (I) or the expression vector; (3) a microarray comprising a CA nucleic acid; (4) an isolated polypeptide encoded comprision of a CA sequence; (5) an isolated antibody, or its antigen binding frame of a CA sequence; (5) an isolated antibody, (6) cc a hybridoma that produces the monoclonal antibody described above; (7) a cc pharmaceutical composition comprising the antibody and a pharmaceutical composition comprising cancer cells, comprising che above (monoclonal) antibody or polypucleotide that selectively comprising cancer cells in an individual; (11) a method for inhibiting growth of cc cancer cells in an individual; (11) a method for delivering a therapeutic agent to cancer cells in an individual; (11) an electronic library comprising the polynucleotide or polypeptide, or their fragments,
   RESULT 123
ADQ59440/c
  S
  Query Match
Best Local S
Matches 46
   carcinoma; (vi) for inhibiting the activity of CAP; (vii) for treating carcinoma; (viii) for neutralizing the effect of CAP; (ix) as a biochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent US2002182586A1, for which no sequence data was published
  New cancer-associated nucleic acid for diagnosing, preventing cancer (e.g. lymphoma) or for screening agents that may be use treating or preventing cancer.
   human; cancer-associated gene; cancer-associated protein; cytostatic; gene therapy; vaccine; tyrosine kinase antagonist; G-protein coupled receptor antagonist; cancer; lymphoma; gene; ds.
   Claim 16;
   Homo sapiens.
  Human cancer-associated (CA)
  ADQ59440;
   ADQ59440 standard; DNA; 347814
   Sequence 276276 BP; 68379 A; 69211 C; 66764 G;
  17-DEC-2002; 2002US-00322696
  15-DEC-2003; 2003WO-US040082
  15-JUL-2004
  WO2004058288-A1
   (SAGR-) SAGRES DISCOVERY INC
  212141 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGG 212186
  2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGG 2933
   2004-543349/52
   DW,
  46;
  Similarity
   SEQ ID NO 76; 143pp; English.
   Malandro
  (first entry)
  1.5%;
   š
  0
  Score 46;
Pred. No.
  gene sequence SEQ ID NO:76
  Mismatches
  DB 11;
. 9e-11;
  0,
   71922 T; 0 U; 0
   Length 276276;
  Indels
  ing or treating used for
  o;
   Other;
  Gaps
  0
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밁 S

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RESULT 124
AAS27638/c
ID AAS27638 standard; DNA; 145 BE
XX
AC AAS27638;
XX

AC AAS27638;
XX

DT 07-NOV-2001 (first entry)
DT 07-NOV-2001 (first entry)
DT 07-NOV-2001 (first entry)
DT 07-NOV-2001 (first entry)
XX

DNA encoding novel signal trar
DE DNA encoding novel signal trar
XX

Neuroprotective; cytostatic; c
XW

Antiinflammatory; anti-HIV; ar
XW

crandiovascular respiratory; r
XW

chromosomal abnormality; Down
XW

Treproductive system; gastroini
XW

Acquired immune deficiency sy:
XX

PO 02-AUG-2001.
XX

PD 02-AUG-2001.
XX

PD 02-AUG-2001.
XX

PD 02-AUG-2001.
XX

PP 04-FEB-2000; 2000US-0179065P.
PR 04-FEB-2000; 2000US-0180628P.
PR 04-FEB-2000; 2000US-0180628P.
PR 04-FEB-2000; 2000US-0180628P.
PR 19-MAR-2000; 2000US-0180628P.
PR 19-MAR-2000; 2000US-0180628P.
PR 11-MAR-2000; 2000US-021886P.
PR 11-MAY-2000; 2000US-021886P.
PR 28-JUN-2000; 2000US-021886P.
PR 28-JUN-2000; 2000US-021886P.
PR 30-JUN-2000; 2000US-021886P.
PR 31-JUL-2000; 2000US-021886P.
PR 31-JUL-2000; 2000US-021886P.
PR 31-JUL-2000; 2000US-021886P.
PR 11-JUL-2000; 2000US-021886P.
PR 11-JUL-2000; 2000US-021889P.
PR 11-JUL-2000; 2000US-021889P.
PR 11-JUL-2000; 2000US-021889P.
PR 11-JUL-2000; 2000US-02189P.
PR 11-JUL-2000; 2000US-02
   mentioned above; (13) a method of screening for anticancer activity; (14) methods for detecting cancer associated with expression of a polypeptide or the presence of the antibody in a test cell or serum sample; (15) a method for screening for a bioactive agent capable of modulating the activity of a CA protein encoded by the above nucleic acid molecule; and (16) a method for treating cancers. (I) has cytostatic activity, and can be used in gene therapy, in vaccines, as a tyrosine kinase antagonist, and as a G-protein coupled receptor antagonist. The compositions and methods of the present invention can be used for diagnosing, preventing and treating cancer, especially lymphomas. They may also be used in compositions and concerning for agents that may be used for treating or preventing cancer. The present sequence represents a human CA gene sequence, which is given in the exemplification of the present invention. Note: The sequence data concerning the sequence concerning concerning cancer contained in electronic format directly from WIPO at the published pot sequences.
  Query Match
Best Local (
   Matches
  Neuroprotective; cytostatic; dermatological; immunosuppressive; tumour; antiinflammatory; anti-HIV; antibacterial; antiinflammatory; cancer; immune system disorder; rheumatoid arthritis; inflammatory condition; organ transplant rejection; infection; hepatitis (; blood disorder; sickle cell anaemis; hyperproliferative disorder; Gaucher's disease; neurodegenerative disorder; Alzhelmer's disease; Parkinson's disease; chromosomal abnormality, Down syndrome; ischaemia; renal disorder; cardiovascular; respiratory; wound healing; endocrine; Addison's disease; reproductive system; gastrointestinal; liver disorder; AIDS; ds;
   Sequence 347814 BP;
   DNA encoding novel signal transduction pathway protein,
   261007
   3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAGAGCTC 3116
  46;
   Similarity
  immune deficiency syndrome.
   CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTC 260962
         2000US-0180628P.
2000US-0186350P.
2000US-0186350P.
2000US-0186350P.
2000US-0198123P.
2000US-0205515P.
2000US-0205467P.
2000US-0215135P.
2000US-0215435P.
2000US-02174880P.
2000US-0217487P.
2000US-0217487P.
2000US-0217487P.
   Conservative
   2001WO-US001312
   2000US-0179065P
   1.5%; but
100.0%; Pr/
   109468 A; 63155 C; 63484 G; 111535 T; 0 U;
  ₽₽
  Score 46;
Pred. No.
   ore 46; DB 12; I red. No. 8.9e-11; Mismatches 0;
   Length 347814;
  Indels
   Seq ID 1298.
  0
  172 Other;
   Gaps
  0
                \forall R \foral
         26-JUII-2000
14-AUG-2000
16-AUG-2000
17-AUG-2000
18-AUG-2000
18-AU
   2000US-0226279P.
2000US-0226681P.
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2000US-0228924P.
2000US-0229343P.
2000US-0229345P.
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2000US-0229343P.
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2000US-0232399P.
  2000US-0220963P.
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2000US-0224519P.
2000US-0225213P.
2000US-0225213P.
2000US-0225266P.
2000US-0225267P.
2000US-0225267P.
2000US-0225267P.
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2000US-0225759P.
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   2000US-0235836P
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   2000US-0236369P
  2000US-02363
  2000US-023
   2000US-0233065P
  2000US-0233064P
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05-DEC-2000
05-DEC-2000
05-DEC-2000
06-DEC-2000
06-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
08-DEC-2000
            The invention relates to novel isolated polypeptides (I), and polynucleotides (II). (II) and the antibody to (I) are useful for diagnosing, preventing and treating diseases including immune system disorders (e.g. congenital and acquired immunodeficiencies, autoimmune disorders (e.g. rheumatoid arthritis), inflammatory conditions, organ transplant rejections and graft versus host disease, infectious diseases (e.g. hepatitis C), bleeding disorders, haemoglobin abnormalities and other blood-related disorders (sickle cell anaemia), myeloproliferative disorders, primary haematopoietic disorders, hyperproliferative disorders (e.g. Gaucher's disease, Parkinson's disease), chromosomal abnormalities (e.g. Gaucher's disease, Parkinson's disease), chromosomal abnormalities (Down syndrome), ischaemic injury (e.g. stroke), renal disorders (e.g.
  Novel polypeptides useful for diagnosing, treating, prognosing disorders related to the proteins, includisorders and neuronal disorders.
   WPI;
(Down syndrome), ischaemic injury (e.g. stroke), renal disorders glomerulonephritis), cardiovascular disorders (e.g. arrhythmia),
  Claim 1;
   Rosen
  17-NOV-2000;
  17-NOV-2000;
17-NOV-2000;
  17-NOV-2000;
17-NOV-2000;
   (HUMA-)
   2001-465460/50.
   ξ
   HUMAN
  SEQ ID NO 1298; 880pp; English.
   Barash SC,
   2000US-0246475P.
2000US-0246476P.
2000US-0246478P.
2000US-0246523P.
2000US-0246523P.
2000US-0246523P.
2000US-0246523P.
2000US-0246523P.
2000US-0246523P.
2000US-0246610P.
2000US-0246611P.
2000US-0249203P.
2000US-0249203P.
2000US-0249213P.
2000US-0249213P.
2000US-0249213P.
2000US-0249213P.
2000US-0249213P.
2000US-0249214P.
2000US-0249213P.
2000US-025921868P.
2000US-0251868P.
2000US-0251989P.
2000US-0251999P.
2000US-025199P.
   GENOME SCI INC.
   Ruben
   ĸ
  including
  preventing and/or
  cancers, immune
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RESULT 125
AAK68506

ID AAK68506

AAK68506

AXX AAK685

AX AAK685

AX AAK685

AX AAK685

AX Human

XX Human

XX Cytost

XX Cytost

XX O9-AUC

PR 17-JAN

XX O9-AUC

X
   ននិនិនិនិនិនិនិ
   밁
  S
  Best Loc
Matches
  Query Match
  14-JUL-2000;
26-JUL-2000;
26-JUL-2000;
14-AUG-2000;
  28-JUN-2000;
30-JUN-2000;
07-JUL-2000;
07-JUL-2000;
11-JUL-2000;
11-JUL-2000;
   02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
17-MAR-2000;
18-APR-2000;
19-MAY-2000;
07-JUN-2000;
  respiratory disorders, dermatological disorders, in wound healing, epithelial cell proliferation, endocrine disorders (e.g. Addison's disease), reproductive system disorders, gastrointestinal disorder (inflammatory disorders), liver disorders, (cirrhosis), as stimulators of B-cell responsiveness to pathogens, activators of T-cells, to induce higher affinity antibodies, and as a means to induce tumour proliferation in pathologies e.g. acquired immune deficiency syndrome (AIDS). AA26976-AAS27850 represent novel signal transduction pathway protein coding sequences and PCR primers of the invention
  31-JAN-2000;
   09-AUG-2001
   cytostatic;
   Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
   Human immune/haematopoietic
  06-NOV-2001
  AAK68506;
  AAK68506 standard; DNA; 145
   Sequence 145
   WO200157182-A2
  Local
   3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGACAGAGACTCTGTCTC 3122
   sapiens.
   120
  1 Similarity
45; Conserv
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                     2000US-018464P

2000US-0189874P

2000US-0199874P

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2000US-0216647P

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2000US-0216647P

2000US-0216880P

2000US-0217487P

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2000US-0217487P

2000US-0217487P

2000US-0217487P

2000US-0217487P

2000US-022513P

2000US-022513P

2000US-0225147P

2000US-0225267P

2000US-0225758P

2000US-0225758P

2000US-0225758P

2000US-0225758P

2000US-0225758P
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  BP;
  (first entry)
  2001WO-US001354.
   2000US-0180628P
  2000US-0179065P
   13
  A; 36 C;
   antigen genomic
   ₽P
  Score 45; DB
Pred. No. 3.4
0; Mismatches
   21
   ç.
   75 T;
  3.4e-10;
hes 0;
  DB 4;
   0
   Ģ
   sequence SEQ
   0 Other;
  de.
  Length 145;
  Indels
   76
  ID NO:23318
  0;
  Сарв
```

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밁
                                Ś
   amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cartivity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by reciffying mutations or deletions in a patient's genome the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the culleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, cancers and treat immune/haematopoietic-related diseases, especially cancers and cancer metatases of haematopoietic antigen genomic to AAK64703 ct cancers from the present invention. AAK54942 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention.
   Query Match
Best Local S
Matches 45
   17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
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17-NOV-2000;
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17-NOV-2000;
10-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
08-DEC-2000;
  Sequence
  Nucleic
  Disclosure;
   Rosen
   AAK54951 to
  (HUMA-) HUMAN
                                      3078
  2001-483426/52.
   Ç,
   26
   ch 1.4%; So 1.4%; So 1.4%; So 1.4%; I similarity 100.0%; I 45; Conservative 0;
  for
   acids encoding human immune/hematopoietic for preventing, diagnosing and/or treating
GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  145
   Barash
   AAK64702
  SEQ ID NO 23318; 3071pp + Sequence Listing; English
   2000US-0249210P

2000US-0249210P

2000US-0249211P

2000US-024924P

2000US-024924P

2000US-024926P

2000US-024926P

2000US-024926P

2000US-024926P

2000US-02592P

2000US-02592P

2000US-0251939P

2000US-0251186P

2000US-0251186P

2000US-0251186P

2000US-0251186P

2000US-025186P

   BP;
  GENOME
  75
   SC,
  >
  SCI
  encode the human immune/haematopoietic antigen (I) given in AAM82170 to AAM91921. (I) have cytostatic
  21
  ü
   Score 45; DB; Pred. No. 3.4
0; Mismatches
  36
   Š
  ຸດ
  13
   DB 4;
3.4e-10;
  Η,
  0
  'n,
  0
  Length 145;
  Other;
  cancers
   Indels
  polypeptides, and metastasis.
   0
   Gaps
```

0,

22-AUG-2000
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23-AUG-2000
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06-SEP-2000
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02-CCT-2000
02-CCT-2000
02-SEP-2000
02-CCT-2000
02-CCT

2000US-0226681P 2000US-0227182P 2000US-022934SP 2000US-022934SP 2000US-022934SP 2000US-022934SP 2000US-02394SP 2000US-023124AP 2000US-023124AP 2000US-023124AP 2000US-023124AP 2000US-023124AP 2000US-023141AP 2000US-023141AP 2000US-023396BP 2000US-023396BP 2000US-023396BP 2000US-023396BP 2000US-02342AP 2000US-02348AP 2000US-02358BP 2000US-02358BP 2000US-0236BP 2000US-0246AP 2000US-0246A

| AND OS-AUG-2001.  AND SOLUTION.  PE 17-JAN-2001 2001W-US001354.  XY 13-JAN-2000 2000US-0149652P.  PR 31-JAN-2000 2000US-0149652P.  PR 44-FEB-2000 2000US-0149654P.  PR 24-FEB-2000 2000US-0149654P.  PR 16-MAR-2000 2000US-014984P.  PR 17-MAR-2000 2000US-014984P.  PR 18-MAG-2000 2000US-014984P.  PR 18-MAG-2000 2000US-025964P.  PR 18-MAG-2000 2000US-025 | RESULT 126 AAK69250 ID AAK69250 standard; DNA; 145 BP.  XX AC AAK69250; XX DT 06-NOV-2001 (first entry) XX DE Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24062. XX KW Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; XX XX OS Homo sapiens. XX PN W0200157182-A2.                                                                                                                                                                                                                                     |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| PR 29-SEP-2000 2000US-024357P. PR 29-SEP-2000 2000US-0233368P. PR 29-SEP-2000 2000US-0233368P. PR 29-SEP-2000 2000US-0233368P. PR 29-SEP-2000 2000US-0233370P. PR 29-CCT-2000 2000US-0237037P. PR 29-CCT-2000 2000US-0237037P. PR 29-CCT-2000 2000US-0237039P. PR 29-CCT-2000 2000US-0239359. PR 29-CCT-2000 2000US-0244950P. PR 29-CCT-2000 2000US-02441786P. PR 29-CCT-2000 2000US-02441787P. PR 29-CCT-2000 2000US-02441786P. PR 29-CCT-2000 2000US-0244617P. PR 29-CCT-2000 2000US-0244477P. PR 29-CCT-2000 2000US-0244478P. PR 29-CCT-2000 2000US-0244611P. PR 29-CCT-2000 2000US-024 | 14-SEP-2000; 2000US-0; 21-SEP-2000; 2000US-0; 21-SEP-2000; 2000US-0; 25-SEP-2000; 2000US-0; 25-SEP-2000; 2000US-0; 26-SEP-2000; 2000US-0; 27-SEP-2000; 2000US-0; 27-SEP-2000; 2000US-0; 27-SEP-2000; 2000US-0; 29-SEP-2000; 2000US-0; |

```
묽
  र्
  Query Match
Best Local Similarity
Matches 45; Conserv
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05-DBC-2000

05-DBC-2000

06-DBC-2000

08-DBC-2000

08-DBC-2000

08-DBC-2000

08-DBC-2000

08-DBC-2000

11-DBC-2000
   amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the nucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-entrice actived cells. AAK64703 to have for the content that the content the content that the content
              acquired immunodeficiency virus; dysphagia; gastrointestinal disorder; adenocarcinoma; reproductive system disorder; testicular feminisation; endocrine disorder; diabetes; cancer; leukaemia; neovascularisation; respiratory disorder; renal disorder; kidney failure; blood disorder; myocardial infarction; wound healing; cell proliferation; skin aging; food additive; food preservative; gene therapy; gene; ds.
  Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis.
  Central nervous system; CNS; autoimmune disease; rheumatoid arthritis; hyperproliferative disorder; neoplasm; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; ischaemia; anglogenesis; nervous system disorder; Alzheimer's disease; AIDS; ocular disorder;
  Genomic
  05-JUN-2002
   ABK44042 standard; DNA; 145 BP
  Sequence 145
  6
  AAK54951
  Disclosure;
   Rosen
   (HUMA-)
   05-JAN
  AAK87694 represent human immune/haematopoietic antigen genomic quences from the present invention. AAK54942 to AAK54950 and AAM82169 present sequences used in the exemplification of the present invention
  3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   Š
  26 GTGCCACTGCAGTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  DNA
   HUMAN GENOME SCI INC.
   to AAK64702 encode the human immune/haemat id sequences given in AAM82170 to AAM91921.
   Barash SC,
   2000US-0251988P.
2000US-0256719P.
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2001US-0259678P.
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  encoding novel central nervous system protein
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100.0%; Pred. No.
  24062; 3071pp + Sequence Listing;
   Ruben
  <u>.</u>
  S
  Mismatches
   immune/haematopoietic antigen (I) to AAM91921. (I) have cytostatic
   3.4e-10;
  DB 4;
  ς;
  0 Other;
  ..
  Length 145;
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  #244.
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19-MAY-2000;
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   17-JAN-2001;
   2000US-0186350P.
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   17-NOV-2000;
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2000US-0251030P.
2000US-0251988P.
2000US-0251479P.
2000US-0251866P.
2000US-0251866P.
2000US-0251869P.
2000US-0251869P.
2000US-0251990P.
2000US-0254997P.
2001US-0259678P.
  2000US-0236802P

2000US-0237038P

2000US-0237039P

2000US-0237039P

2000US-023993P

2000US-0241921P

2000US-0241786P

2000US-0241808P

2000US-0241808P

2000US-0246474P

2000US-0246474P

2000US-0246477P

2000US-0246477P

2000US-0246477P

2000US-0246477P

2000US-0246477P

2000US-0246524P

2000US-0246524P

2000US-0246524P

2000US-0246524P

2000US-0246524P

2000US-0246521P

2000US-0246521P

2000US-024651P

2000US-024651P

2000US-024651P

2000US-024651P

2000US-024651P

2000US-024921P

  2000US-0235834P.
2000US-0235836P.
2000US-023632P.
2000US-0236367P.
2000US-0236369P.
2000US-0236369P.
2000US-0236369P.
2000US-0236369P.
  GENOME SCI
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l Similarity
45; Conserv

Conservative

0;

Mismatches

0,

Indels

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Gaps

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3122

100.0%;

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Query Match
Best Local S
Matches 45
  leukaemia, disorders involving neovascularisation e.g. malignancies, respiratory disorders e.g. nonallergic rhinitis, renal disorders e.g. acute kidney failure and blood related disorders e.g. myocardial infarction. The polypeptides can also be used to aid wound healing and epithelial cell proliferation, to prevent skin aging due to sunburn, to maintain organs before transplantation, for supporting cell culture of primary tissues, to regenerate tissues and in chemotaxis. The
  e.g. corneal infection, gastrointestinal disorders e.g. dysphagia, adenocarcinomas and irritable bowel syndrome, reproductive system disorders e.g. testicular feminisation, endocrine disorders e.g. diabete and pituitary dwarfism, cancers and disorders at the cellular level e.g.
   angiogenesis, nervous system disorders e.g. Alzheimer's disease and amylotrophic lateral sclerosis, infections caused by bacteria, viruses e.g. Acquired immunodeficiency virus (AIDS) and fungi, ocular disorders
  carbohydrate,
  polypeptides can also be used as a food additive or preservative to increase or decrease storage capabilities, fat content, lipid, prote
  disorders e.g. neoplasms of the breast or liver, cardiovascular disorde e.g. cardiac arrest, cerebrovascular disorders e.g. cerebral ischaemia,
   pathological condition. Disorders which are diagnosed or treated include autoimmune diseases e.g. rheumatoid arthritis, hyperproliferative disorders e.g. neoplasms of the breast or liver, cardiovascular disorders
   Disclosure; SEQ ID NO 1230;
   New isolated nucleic acid encoding a protein for diagnosing, treating or ameliorating medical conditions and used as food
   Rosen
  novel
   E)
   2001-581633/65
   central
  δ
  ention describes an isolated nucleic acid molecule (I) encoding a entral nervous system protein. (I) and polypeptides (III) encoded are used to treat a medical conditions and in diagnosis of a
   Barash
  vitamins,
  SC,
  1.4%;
   Ruben
   minerals,
  837pp; English.
                       Score 45;
Pred. No.
   XS.
  cofactors and other nutritional
                       DB 4; Lo
  Length 145
  preventing, additives c
   protein,
   diabetes
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RESULT 128
ADB94441/c
ID ADB944
XX Increa
XX Homo s
XX Homo s
XX Increa
XX I
   밁
   Ś
31-JAN-2000;
04-FEB-2000;
28-JUN-2000;
07-JUL-2000;
   ds; gene; human; autoimmune disease; Parkinson's disease; silicosis; gastrointestinal disease; atherosclerosis; haemophilia; thrombocytopenia; immunosuppressive agent; adjuvant; enhance immune response; higher affinity antibody induction; increased serum immunoglobulin concentration.
  04-DEC-2003
   ADB94441
   17-JAN-2001;
   US2002168711-A1
   Homo sapiens.
  Novel human protein DNA #50.
  ADB94441;
  14-NOV-2002.
   26 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 70
   standard; DNA; 145
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
2000US-0179065P.
2000US-0180628P.
2000US-0214886P.
2000US-0216647P.
   2001US-00764868.
  (first entry)
   ВP
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11-JUL-2000

11-JUL-2000

11-JUL-2000

26-JUL-2000

26-JUL-2000

14-AUG-2000

14-AUG-2000

14-AUG-2000

14-AUG-2000

14-AUG-2000

14-AUG-2000

14-AUG-2000

14-AUG-2000

10-SEP-2000

22-AUG-2000

01-SEP-2000

01-SEP-2000

01-SEP-2000

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01-SEP-2000

02-SEP-2000

21-SEP-2000

22-SEP-2000

29-SEP-2000

29-SEP-2000

29-SEP-2000

29-SEP-2000

29-SEP-2000

29-SEP-2000

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21-SEP-2000

21-SE
            The invention relates to an isolated polypeptide. The polypeptide is useful for diagnosing a pathological condition or a susceptibility to a pathological condition in a subject, by determining the presence or amount of expression of the polypeptide in a biological sample and diagnosing a pathological condition or a susceptibility to a pathological condition or a susceptibility to a pathological condition based on the presence or amount of expression of the polypeptide is also useful for identifying a binding partner to the polypeptide is also useful for identifying a binding partner to the polypeptide, which involves contacting the polypeptide with a binding partner and determining whether the binding partner effects an activity of the polypeptide. The polypeptide or the nucleic acid encoding the polypeptide is useful for preventing, treating, or amendment of the contacting the polypeptide is useful for preventing.
  08-DEC-
  (ROSE/)
(RUBE/)
(BARA/)
   Rosen
ameliorating a medical
  New isolated polypeptide useful for diagnosing and treating
   ξ
  ROSEN C
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   Ruben SM,
  2000US-021680P

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2000US-0217496P

2000US-0220964P

2000US-022968P

2000US-0225267P

2000US-0225267P

2000US-022547P

2000US-022547P

2000US-022548P

2000US-022548P

2000US-022548P

2000US-0225757P

2000US-0225868P

2000US-0225889

2000US-022934P

2000US-022934P

2000US-022934P

2000US-022934P

2000US-023934P

2000US-023934P

2000US-0239389

2000US-0234274P

2000US-0234274P

2000US-0234274P

2000US-0234274P

2000US-023668P

2000US-023668P

2000US-0237039P

2000US-0237039P

2000US-0237039P

2000US-0237039P

2000US-0237039P

2000US-0241785P

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2000US-024461PP

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2000US-0251868P

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  SEQ ID NO
  က က
  ß
  3 Þ
  1298;
   Barash
  condition,
  345pp;
   SC
  which
  English.
  involves
  administering
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Parkinson's

31-JAN-2000; 04-FBB-2000; 24-FBB-2000; 02-MAR-2000; 16-MAR-2000; 17-MAR-2000; 18-APR-2000;

2000US-0179065P. 2000US-0180628P. 2000US-0184664P. 2000US-0186350P. 2000US-0189874P. 2000US-0199076P. 2000US-0199075P.

17-JAN-2001; 29-JAN-2004

2001US-00764875

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ARESULT 129
ANDISSO
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ADISSO
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  cc is useful for diagnosing a pathological condition or a succeptibility to cc a pathological condition in a subject, which involves determining the cc presence or absence of a mutation in The nucleic acid, and diagnosing a compatibility to a pathological condition or susceptibility to a pathological condition or cc based on the presence or absence of the mutation. The polypeptide, the cc nucleic acid and an antibody to the polypeptide are useful for treating cc autoimmune disease, Parkinson's disease, silicosis, gastrointestinal cc disease, atherosclerosis, haemophilia, thrombocytopenia. The polypeptide, cc the nucleic acid and the antibody are useful as immunosuppressive agents, cas adjuvants to enhance immune responses, and as agents to induce higher cc affinity antibodies and increase serum immunoglobulin concentrations. The cr present sequence represents DNA encoding a novel human protein. Note: The sequence data for this patent did not form part of the printed concentrations by a specification but was obtained in electronic format direct from USPTO at cardial nearty convisions.
   Best Loc
Matches
  Query Match
  immune system disorder; diabetes; rheumatoid arthritis; isystemic lupus erythematosus; autoimmune thyroiditis; haemolytic anaemia; systemic lupus erythematosus; autoimmune thyroiditis; haemolytic anaemia; inflammatory disorder; ischaemia-reperfusion injury; inflammatory bowel disease; Crohn's disease; infectious disease; HIV infection; hepatitis infection; bacterial infection; fungal infection; parasitic infection; muscular disorder; reproductive disorder; gastrointestinal disorder; pulmonary disorder; cardiovascular disorder; atherosclerosis; arrhythmia; myocarditis; renal disorder; acute glomerulonephritis; pyelonephritis; pyelonephritis; pyolonephritis; pyolonephritis; pyolonephritis; human; ds.
  polypeptide or the nucleic acid to a mammalian subject. The nucleic acid is useful for diagnosing a pathological condition or a susceptibility to a pathological condition in a subject, which involves determining the presence or absence of a mutation in The nucleic acid, and diagnosing a
  anti-HIV; hepatotropic; virucide; antihyroid; anti-anemic; vasotropic; anti-HIV; hepatotropic; virucide; antibacteria; fungicide; antiparasitic; muscular; gynaecological; gastrointestinal; respiratory; cardiovascular; antiarteriosclerotic; antiarrhythmic; cardiant; nephrotropic; litholytic; cytostatic; gene therapy; neural disorder; Alzheimer; disease; Parkinson's disease; Huntington's chorea; amyotrophic lateral sclerosis; multiple sclerosis;
   neuroprotective; nootropic; antiparkinsonian; anticonvulsant; antidiabetic; antirheumatic; antiarthritic; dermatological;
  22-APR-2004
  ADI55027
  US2004018969-A1.
   ADI55027;
  seqdata.uspto.gov/sequence.html?DocID=20020168711.
  Local
  3078
   120
  human
   l Similarity
45; Conserv
   145
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   standard;
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   1.4%;
ilarity 100.0%;
Conservative
   protein genomic
  BP; 13
  (first
   DNA;
  entry)
   A; 36
   145
   Ç
   0;
  DNA seq id 1230.
   В₽
  Score 45;
Pred. No.
   21
   <u>.</u>
  Mismatches
  75 T;
   ŏ.
  DB 10;
3.4e-10
   0 U; 0
  0
  Length
   Other;
  Indels
   3122
   76
  0,
   Gaps
  0
```

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20-OCT-2000

20-OCT-2000

20-OCT-2000

20-OCT-2000

20-OCT-2000

01-NOV-2000

08-NOV-2000

17-NOV-2000

17-NO
   New polypeptides and nucleic acid molecules, useful for detecting, preventing, diagnosing, prognosticating, treating or ameliorating medic conditions e.g. neural disorders, reproductive disorders or infectious diseases.
  (ROSE/)
(RUBE/)
(BARA/)
                                Disclosure;
  2004-122079/12.
   Š
  ROSEN C
RUBEN S
BARASH
   Ruben
                                SEQ
   2000US-0241221P-
2000US-0241785P-
2000US-0241809P-
2000US-0246475P-
2000US-0246475P-
2000US-0246477P-
2000US-0246477P-
2000US-0246478P-
2000US-0246478P-
2000US-0246528P-
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2000US-0251989P-
2000US-0251989P-
2000US-0251999P-
2000US-0251990P-
2000US-0251990P-
2000US-0251990P-
2000US-0259678P-
  CA.
SM.
                                IJ
   SM,
                              NO 1230; 413pp; English
   Barash
   SC
   medical
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The

invention describes an isolated polypeptide comprising

an

amino acid

19-MAY-2000
28-JUN-2000
28-JUN-2000
28-JUN-2000
29-JUN-2000
11-JUL-2000
11-JUL-2000
11-JUL-2000
14-JUL-2000
14-JUL

20000US.

S.-020545P S.-02154886P S.-0216486P S.-0216480P S.-0216480P S.-0225963P S.-02252513P S.-02252513P S.-022525759P S.-0225758P S.-0225759P S.-0225759P S.-0225759P S.-0225759P S.-0225759P S.-0226681P S.-0227009P S.-0227009P S.-0229344P S.-02314413P S.-0231444P S.-02314413P S.-02314413P S.-02314413P S.-023163P S.-0231639P 
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RESULT 130
ACH37117
ACH3711
XX
ACH3711
XX
ACH371
ACH3
   Query Match
Best Local S
Matches 45
The invention relates to an isolated polynucleotide comprising any one o 38043 cDNA sequences, appearing as ACH12789-ACH50831, whose sequence was determined by the technique of SBH (sequencing by hybridisation). Also included is a purified polypeptide comprising a sequence corresponding t a reading frame of the novel polynucleotide. The nucleic acid sequences are useful in diagnostics as expressed sequence tags (EST) for identifying expressed genes or for physical mapping of the human genome, in forensics, in assessing biodiversities, or in identifying mutations responsible for genetic disorders and other traits. The nucleotide sequences are also useful as hybridisation probes, as oligomers for PCR, for chromosome and gene mapping, in the recombinant production of protein, or in generating antisense DNA or RNA. The purified polypeptide
  sequence at least 90% identical to: a polypeptide fragment, domain, epitope, or full-length protein of any one of 607 amino acid sequences (I) described in the specification; a polypeptide fragment of (I), or the encoded sequence contained in (II), having biological activity; or a variant, allelic variant, or a species homologue of (I). The polypeptides and nucleic acid molecules are useful for detecting, preventing, diagnosing, prognosticating, treating or ameliorating medical conditions such as neural disorders, e.g. Alzheimer's disease, Parkinson's disease, Huntington's chorea, amyotrophic lateral sclerosis or multiple sclerosis, immune system disorders, e.g. diabetes, Theumatoid arthritis, systemic
   New polynucleotide sequences obtained as hybridization probes, as oligomers mapping, in the recombinant production
  genome
   (LABA/)
(STAC/)
(DICK/)
   Human endothelial cell cDNA #5250
  ACH37117;
   ACH37117 standard; cDNA;
   30-JUL-2001; 2001US-00918995
  30-JUL-2001; 2001US-00918995
   17-APR-2003.
   US2003073623-A1
   13-OCT-2003
  (DRMA/)
  3078
   2003-615964/58.
   sapiens
   erythematosus,
  26
  ss; sequencing
mapping; biodiv
   DICKSON M C.
JONES L W.
   45
   RT,
   Similarity
   LABAT I.
  DRMANAC R T.
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 70
  DNA or RNA.
   Conservative
   ID NO 24329; 44pp;
   Labat I,
  (first entry)
  biodiversity;
   1.4%;
  autoimmune thyroiditis or haemolytic anaemia,
   w
   by hybridisation; SBH; expressed sequence tag; EST; versity; genetic disorder.
  Stache-Crain B,
  492
   0;
  Score 45;
Pred. No.
   ВP
  English
   Mismatches
   n of protein,
  3.4e-10;
hes 0;
  В
  Dickson MC,
   12;
   Length 145;
   Indels
  cDNA libraries, useful chromosome and gene or in generating
  Jones
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   0
   Gaps
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   0
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RESULT 131
ANAH13294/c
ID ANAH132
XX ANH132
XX ANH132
XX Human
DE Human
XX Human;
XX EP1074
XX O7-FEE
PM 29-JUI
PM 29-JUI
PM 27-AUC
PM 01-JAUC
PM 09-JUN
XX HELI-
XX Ota T,
PI Ishii
XX PF CLaim
XX WPI;
XX PF CLaim
XX PF CLaim
XX PF CLaim
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XX PF CC CAIL
YC CLAIM
XX PF CC CAIL
YC  អន្តិ និង្ស
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   S
                         The present invention describes primer sets for synthesising 5602 full-
CC length cDNAs defined in the specification. Where a primer set comprises:
CC (a) an oligo-cT primer and an oligonucleotide complementary to the
CC complementary strand of a polynucleotide which comprises one of the 5602
CC conclectide sequences defined in the specification, where the
CC coligonucleotide comprises at least 15 nucleotides; or (b) a combination
CC oligonucleotide comprises at least 15 nucleotides; or (b) a combination
CC of an oligonucleotide comprising a sequence complementary to the
CC complementary strand of a polynucleotide which comprises a 5'-end
CC sequence and an oligonucleotide comprising a sequence complementary to a
CC polynucleotide which comprises a 3'-end sequence, where the
CC oligonucleotide which comprises a 1 least 15 nucleotides and the combination of
CC the 5'-end sequence'3'-end sequence is selected from those defined in the
CC specification. The primers are useful for synthesising polynucleotides,
CC particularly full-length CDNAs. The primers are also useful for the
CC particularly full-length CDNAs. The primers are also useful for the
  Matches
  Query Match
Best Local
   29-JUL-1999;
27-AUG-1999;
11-JAN-2000;
02-MAY-2000;
09-JUN-2000;
   is useful for generating antibodies specific for it. is one of the 38043 isolated cDNA/SST sequences. Note for this patent did not form part of the printed spec obtained in electronic format directly from USPTO at
   Primer sets for synthesizing polynucleotides, length cDNAs defined in the specification, an diagnosis of the abnormality of the proteins
   EP1074617-A2
   26-JUN-2001
   AAH13294
   Sequence 492 BP; 121 A; 128 C; 137 G; 106 T; 0
  seqdata.uspto.gov/sequence.html?DocID=20030073623
  WPI; 2001-318749/34
  28-JUL-2000; 2000EP-00116126.
   07-FEB-2001.
   2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG
   sapiens
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   တ
   cDNA clone (3'-primer) SEQ ID NO:10129.
   1 Similarity
   primer;
  HELIX RES INST
  Isogai T, Nishikawa T,
Sugiyama T, Wakamatsu
   SEQ
  standard; cDNA;
  TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG
  ; 2000JP-00118776.
; 2000JP-00183767.
; 2000JP-00241899.
  Conservative
  Ħ
   (first
  detection; diagnosis; antisense
  99JP-00248036.
99JP-00300253.
   NO 10129;
   1.4%;
   entry)
   568
   2537pp + Sequence Listing; English
   0
  Score 45;
Pred. No.
   Mismatches
  Hayashi K,
A, Nagai K,
   3 B
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  .3e-10;
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   Saito K,
C, Otsuki
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  encoded
   particularly the 56
d for the detection
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   therapy;
  Length 492
   U; 0 Other;
   Note: The seques specification,
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  410
   Yamamoto
  gene therapy;
   <u>.</u>
   Gaps
  but was
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detection and/or diagnosis of the

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AEB33439
XX AEB3
AC AEB3
AC AEB3
XX OB-6
XX SNP
XX Homc
XX Homc
XX Home
XX H
   RESULT 132
                       The invention relates to a detection reagent capable of detecting one or compressing the nucleic acid polymorphisms. The invention also relates to compute the sub-region, a computer readable medium having stored in it the SNP called molecule for detecting at least specification, an isolated nucleic acid molecule for detecting at least one SNP given in the specification comprising at least about 12 contiguous nucleotides, genotyping at least come SNP position given in the specification in a sample, identifying an comprising at least one container containing a disorder and a kit comprising at least one container containing the detection reagent.

CC Determining whether a trait is linked to one of the human chromosomes or containing whether the trait is linked to one container containing whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises determining whether the trait is linked to one comprises contacting the comprises contacting the comprise with a detection reagent that differentiates between alternative all least one contaction in a sample comprise contaction in a sample comprise contaction in a sample comprise contaction the contaction in the contaction contaction in a sample comprise contaction in a sample comprise contaction in a sa
   Best Loc
Matches
   Naik
  the full-length cDNAs. The primers allow obtaining of the full-length cDNAs easily without any specialised methods. AAH33166 to AAH33628 and AAH33632 to AAH3642 represent human achases sequences; AAB92446 to AAB95893 represent human amino acid sequences; and AAH33629 to AAH3362 represent oligomucleotides, all of which are used in the exemplification of the
  New detection reagent capable of detecting 1, 100, 500, 1000 or 5000 or more single nucleic acid polymorphisms, useful in identifying an individual having or at risk of developing type II diabetes or obesity.
   WPI; 2005-511776/52
   08-SEP-2000;
   07-JUL-2005
   US2005147987-A1
  Homo
   metabolic
   Human DNA
   08-SEP-2005
   Sequence 568 BP; 139
  10-SEP-2001;
   19-JUL-2004; 2004US-00893315.
  antidiabetic;
   AEB33439
   AEB33439 standard;
  (APPL-)
  Local
   detection;
  ,
A
  3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  ds.
   sapiens.
  13;
  47
  APPLERA CORP NY
   invention
  Similarity
   Subramanian
  cic; anorectic; endocrine disease; gastrointestinal disease;
disorder; nutritional disorder; single nucleotide polymorphism;
   polymorphic
  Zhang JN,
   Conservative
  2000US-0231397P.
2001US-00948947.
   (first entry)
  ij
  ; diagnosis; non-insulin dependent diabetes; obesity; anorectic; endocrine disease; gastrointestinal disea
  NO 1202; 31pp;
   DNA;
   တ
   A; 144 C; 119
  100.0%;
   region #1019.
   1.4%;
  Liu X,
   601
   tu X, Rowe W,
Woodage T;
position given in the specification,
   BP
  Score 45; DB; Pred. No. 3.3
  0,
  English
   G; 163 T; 0 U; 3 Other;
   DB 4; Lo
  Cravchik A,
  0,
  Length 568
   Indels
  Kalush
  ম
  ç
  Gaps
  0
```

```
RESULT 133
AAK63029
 밁
  Ś
   នននិន្នននិន្ននិង
   Matches
  Query Match
Best Local
   26-JUL-2000;
26-JUL-2000;
14-AUG-2000;
  determining which allele is present at the at least one SNP position. Identifying an individual having or at risk of developing a disorder comprises genotyping at least one SNP given in the specification in a nucleic acid sample from the individual. The disorder is type II diabetes (non-insulin dependent diabetes) or obesity. The detection reagent is useful in identifying an individual having or at risk of developing a disorder, particularly type II diabetes or obesity. This sequence represents a human DNA polymorphic region used in the scope of the invention. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format from
  14-AUG-2000;
14-AUG-2000;
   11-JUL-2000;
  07-JUL-2000
  30-JUN-2000;
   19-MAY-2000;
07-JUN-2000;
  17-MAR-2000;
18-APR-2000;
   Homo sapiens.
  Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis; ss.
  Human immune/haematopoietic antigen encoding cDNA SEQ ID NO:8089.
  06-NOV-2001
  AAK63029;
   AAK63029 standard; cDNA; 1664
  Sequence
  14-AUG-2000;
  11-JUL-2000;
   17-JAN-2001;
  WO200157182-A2
  USPTO
   2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACA 2939
   202 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACA 246
  at seqdata.uspto.gov/sequence.html.
   45;
  Similarity
   601
2000US-0186350P.
2000US-0199076P.
2000US-0199076P.
2000US-0199076P.
2000US-0216486P.
2000US-021648P.
2000US-021648P.
2000US-021748P.
2000US-021748P.
2000US-021748P.
2000US-021748P.
2000US-021748P.
2000US-021748P.
2000US-02251829P.
2000US-0225113P.
2000US-0225113P.
2000US-0225266P.
2000US-0225266P.
2000US-0225247P.
2000US-0225244P.
2000US-0225244P.
2000US-0225758P.
   Conservative (
   (first entry)
   BP; 174 A; 133 C; 164 G; 129 T; 0 U; 1 Other;
   2001WO-US001354
  1.4%;
   0;
   Score 45;
Pred. No.
   ₽P
   Mismatches
   DB 14;
  3.3e-10;
  0;
  Length 601;
  Indels
   0;
  Gaps
```

```
á
   CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cartivity, and can be used in gene therapy and vaccine production. (I) coroteins and polymucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For cexample, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome capture the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) complete acids into a host cell and culturing the cell to express the used to produce the secreted (I), by inserting the culeic acids into a host cell and culturing the cell to express the cancer and cancer metastases of haematopoietic-related diseases, especially concers and cancer metastases of haematopoietic antigen genomic to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK54942 to AAK5490 and AAM82169
  Query Match
Best Local S
Matches 45
  08-NOV-2000
17-NOV-2000
17-NOV
   Sequence 1664 BP;
   Claim
   Nucleic acids encoding useful for preventing,
  Rosen
  (HUMA-)
2898 GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
   2001-483426/52.
DB; AAM90248.
   Ç,
   45; Conserv
  HUMAN
   SEQ ID NO 8089;
  sequences
  Barash
  2000US-0246611P
2000US-0246613P
2000US-024920PP
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249211P
2000US-0249218P
2000US-0249218P
2000US-0249264P
2000US-0249264P
2000US-0249265P
2000US-0249265P
2000US-0249269P
2000US-0251989P
2000US-02511866P
2000US-02511869P
2000US-0251989P
  1.4%;
ilarity 100.0%;
Conservative
  GENOME
   549
  sc,
  used
  SCI
   ?
  human immune/hematopoietic antigen polypeptides, diagnosing and/or treating cancers and metastasis.
   335
  in the
   3071pp + Sequence Listing; English.
   0,
  Score 45;
Pred. No.
   Ç
  Š
   391
  exemplification
   Mismatches
   ç,
   376 T;
   .2e-10;
  4.
   0 U;
  Length 1664;
  of the
   Indels
   13
   Other;
  present invention
   0
   Gaps
```

114-AUG-2000
22-AUG-2000
22-AUG-2000
22-AUG-2000
23-AUG-2000
21-SEP-2000
01-SEP-2000
01-SEP-2000
01-SEP-2000
06-SEP-2000
06-SEP-2000
08-SEP-2000
08-SEP-2000
08-SEP-2000
08-SEP-2000
08-SEP-2000
114-SEP-2000
115-CCT-2000
115-CCT

2000US-0225759P.
2000US-022688P.
2000US-022688P.
2000US-022688P.
2000US-02270892P.
2000US-0229343P.
2000US-0229343P.
2000US-0229343P.
2000US-0229513P.
2000US-0231243P.
2000US-0231414P.
2000US-0232398P.
2000US-0232398P.
2000US-023493P.
2000US-0244679P.
2000US-0246479P.
2000US-0246479P.
2000US-0246528P.
2000US-0246528P.
2000US-0246538P.
2000US-0246539P.
2000US-0246609P.
2000US-0246609P.

| PR 08-SEP-2000 PR 14-SEP-2000 PR 25-SEP-2000 PR 25-SEP-2000 PR 27-SEP-2000 PR 27-SEP-2000 PR 29-SEP-2000 PR 29- |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |

```
AAK69885
ID AAK698
XX AAK698
XX AAK698
XX D6-NOV
XX Human
XX Human
XX Cytost
XX Cytost
XX Homo s
XX Homo s
XX Homo s
XX Homo s
XX PN WO2001
XX PD 09-AUC
  밁
   S
  amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cacivity, and can be used in gene therapy and vaccine production. (I) cyrotesins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For certainsent of diseases associated with inappropriate (I) expression. For certainsent of diseases associated with inappropriate (I) expression. For certainsent of diseases associated with decreased certainsent of the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the current acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, and cancer metastases of haematopoietic-derived cells. AAK64703 cells. AAK64703 concers and cancer metastases of haematopoietic antigen genomic cancers from the present invention. AAK54942 to AAK54950 and AAM82169 crepresent sequences used in the exemplification of the present invention.
   Query Match
Best Local S
   Matches
  17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
06-DEC-2000;
08-DEC-2000;
   Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis.
   Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis; ds.
  Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24697.
  Sequence 2219
   Rosen
    09-AUG-2001.
                            WO200157182-A2
  Homo sapiens.
  06-NOV-2001
  AAK69885;
   AAK69885 standard; DNA;
   AAK54951
   Disclosure;
   WPI; 2001-483426/52.
   (HUMA-)
   acid e
   ξ
   45;
   HUMAN
  Similarity
   Barash SC,
  1.4%; Sccilarity 100.0%; Pr
Conservative 0;
  2000US-0249300P.
2000US-025031PP.
2000US-0251030P.
2000US-0251988P.
2000US-0251479P.
2000US-0251856P.
2000US-0251868P.
2000US-0251869P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0251989P.
2000US-0259678P.
  AAK64702 encode the human immune/haematopoietic antigen (I) sequences given in AAM82170 to AAM91921. (I) have cytostatic
   SEQ ID NO
  (first entry)
  BP; 633
   GENOME SCI INC
  A; 508 C; 519 G;
   24698; 3071pp + Sequence Listing;
   Ruben
   2219
  Score 45;
Pred. No.
   X.
   ВP
   Mismatches
  DB 4; Lear
3.1e-10;
   559 T;
  0 U; 0 Other;
  Length 2219;
   Indels
  English.
   0
   Gaps
   0
    14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
14-AUG-2000
12-AUG-2000
22-AUG-2000
22-AUG-2000
23-AUG-2000
01-SEP-2000
  31-JAN-2000;
04-FEB-2000;
24-FEB-2000;
02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
18-APR-2000;
  28-JUN-2000;
30-JUN-2000;
07-JUL-2000;
07-JUL-2000;
   19
   17-JAN-2001;
  2-MAR-2000;
5-MAR-2000;
7-MAR-2000;
8-APR-2000;
9-MAY-2000;
   -JUN-2000,
  2000US-018464P.
2000US-019875P.
2000US-0199076P.
2000US-0199076P.
2000US-0199076P.
2000US-0215115P.
2000US-0215115P.
2000US-0215115P.
2000US-0215115P.
2000US-0217487P.
2000US-0217487P.
2000US-0217487P.
2000US-0225213P.
2000US-0225213P.
2000US-0225213P.
2000US-0225213P.
2000US-02252147P.
2000US-02252147P.
2000US-0225267P.
2000US-02257579P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-0225758P.
2000US-023144P.
2000US-023144P.
2000US-0231244P.
2000US-0231249P.
2000US-0233499P.
2000US-0233499P.
2000US-023484P.
2000US-0235834P.
2000US-0235834P.
2000US-0235836P.
2000US-0236367P.
2000US-0236367P.
2000US-0236367P.
   2001WO-US001354
```

```
39-SEP-2000, 2000US-0216599.
39-SEP-2000, 2000US-0216599.
39-CCT-2000, 2000US-0216599.
39-CCT-2000, 2000US-021690.
31-CCT-2000, 2000US-021690.
```

```
RESULT 136
AAK86256/c
  밁
  S
  cc amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cc activity, and can be used in gene therapy and vaccine production. (I) converted to treatment of diseases associated with inappropriate (I) expression. For cc example, they may be used to treat disorders associated with decreased cc expression by rectifying mutations or deletions in a patient's genome cc that affect the activity of (I) by expressing inactive proteins or to cc supplement the patients own production of (I). Additionally, (I) cc polynucleotides may be used to produce the secreted (I), by inserting the cc protein. (I) proteins and polynucleotides may be used to produce the secreted (I), by inserting the cc protein. (I) proteins and polynucleotides may be used to prevent, cc diagnose and treat immune/haematopoietic related diseases, especially cc cancers and cancer metastases of haematopoietic actived cells. AAK64703 cc sequences from the present invention. AAK54942 to AAK54950 and AAM82169 crepresent sequences used in the exemplification of the present invention.
  Best Loc
Matches
  Query Match
             31-JAN-2000
04-FEB-2000
24-FEB-2000
12-MAR-2000
11-MAR-2000
11-MAR-2000
11-MAR-2000
07-JUN-2000
07-JUN-2000
07-JUN-2000
07-JUL-2000
07-JUL-2000
11-JUL-2000
  WO200157182-A2
   cytostatic;
  Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
   Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41068
   07-NOV-2001
  AAK86256;
  AAK86256
  Sequence 2219 BP;
   Disclosure; SEQ ID NO 24697; 3071pp + Sequence Listing; English
  Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis.
  17-JAN-2001;
  09-AUG-2001.
   Local
   1966 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 2010
  2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG
  1 Similarity
45; Conserv
  standard; DNA; 2219
  to AAK64702 encode the human
   gene
2000US-0186350P.
2000US-019874P.
2000US-0199076P.
2000US-0198123P.
2000US-0205515P.
2000US-020148867P.
2000US-02148867P.
2000US-0216847P.
2000US-0216847P.
2000US-0217486P.
2000US-0217496P.
2000US-0217496P.
2000US-0217496P.
2000US-0220964P.
  1.4%; Score 45; DB 4; larity 100.0%; Pred. No. 3.1e-Conservative 0; Mismatches
   2001WO-US001354.
   (first entry)
   2000US-0180628P.
2000US-0184664P.
   therapy; vaccine; metastasis;
   633 A; 508 C;
  BP
  519 G;
   immune/haematopoietic antigen (I) to AAM91921 (I) have cytostatic
  559 T; 0 U; 0 Other;
   3.1e-10;
   <u>;</u>
  Length 2219;
  Indels
   2932
   0
   Gaps
   0
```

```
CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic cactivity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased cc expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I), by inserting the CC protein. (I) proteins and polynucleotides may be used to produce the secreted (I), by inserting the CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic actived cells. AAK64703 to AAK87694 represent human immune/haematopoietic antigen genomic CC sequences from the present invention. AAK54942 to AAK54950 and AAM82169
   08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
08.NOV-2000
01.7.NOV-2000
17.NOV-2000
17.N
   Nucleic
  Rosen
   Disclosure;
   useful
   (HUMA-)
   2001-483426/52.
  Ş
   c acids encoding for preventing,
   HUMAN
  Barash
   SEQ
   2000US-0246477P
2000US-0246477P
2000US-0246523P
2000US-0246528P
2000US-0246532P
2000US-0246532P
2000US-0246611P
2000US-0246611P
2000US-0246611P
2000US-0249211P
2000US-0249219P
2000US-0249219P
2000US-0249219P
2000US-0249219P
2000US-0249219P
2000US-0249219P
2000US-02592186P
2000US-0259319P
2000US-02511886P
2000US-02511869P
   GENOME
   Ħ
  sc,
   ö
   SCI
   41068; 3071pp +
  human immune/hematopoietic diagnosing and/or treating
  Ruben
  S
   Sequence Listing;
   antigen polypeptides, cancers and metastasis.
```

English.

the

and

14-AUG-2000
12-AUG-2000
22-AUG-2000
23-AUG-2000
01-SEP-2000
02-OCT-2000
03-NOV-2000
08-NOV-2000

2000US-0224518P
2000US-0225214P
2000US-0225214P
2000US-0225268P
2000US-0225268P
2000US-022575PP
2000US-0231043PP
2000US-0231243PP
2000US-0231239PP
2000US-0231243PP
2000US-02441829PP
2000US-02446474PP
2000US-0246474PP
2000US-0246474PP

17-NOV-2000; 17-NOV-2000; 17-NOV-2000;

17-NOV-2000;

```
RESULT 138
ADR07058
ID ADR070
XX
AC ADR070
XX
DT 04-NOV
XX
  S
  밁
   ARKS4951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic CC activity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC encleic acids into a host cell and culturing the cell to express the CC protein. (I) proteins and polynucleotides may be used to prevent, CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic actived cells. AAK64703 CC to AAK87694 represent human immune/haematopoietic antigen genomic complete sequences from the present invention. AAK54942 to AAK54950 and AAM82169 crepresent sequences used in the exemplification of the present invention of the present invention.
  Matches
   Query Match
Best Local Similarity
  17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
01-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
06-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
   Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastass
   17-NOV-2000;
17-NOV-2000;
                04-NOV-2004
  Disclosure; SEQ ID NO 41069; 3071pp + Sequence Listing; English
   ADR07058 standard;
  Sequence 2219 BP; 559 A; 519 C; 508 G; 633 T;
   WPI; 2001-483426/52.
   Rosen
  ADR07058
  (HUMA-) HUMAN GENOME
   2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 2932
  254
   ξ
  45;
   Barash SC,
   2000US-0250391P.
2000US-0251030P.
2000US-0251980P.
2000US-0256719P.
2000US-0251879P.
2000US-0251868P.
2000US-0251869P.
2000US-0251869P.
2000US-0251869P.
  2000US-0249215P.
2000US-0249216P.
2000US-0249218P.
2000US-0249218P.
2000US-0249244P.
2000US-0249265P.
2000US-0249265P.
2000US-0249269P.
2000US-024929P.
2000US-024929P.
2000US-024929P.
2000US-024929P.
2000US-024920P.
   Conservative
                (first entry)
   CDNA; 2491
   1.4%;
  SCI INC
   Ruben
   Score 45; pred. No.
   0,
   X.
   ВP
  Mismatches
   DB 4;
   3.1e-10;
  0 U; 0 Other;
  0
  Length 2219;
  Indels
  and metastasis
   0
   Gaps
   0
```

```
RESULT 139
AAH18230
ID AAH182
XX
AC AAH182
XX
DT 26-JUN
XX
DB Human
  S
   닭
  Query Match
Best Local (
   Matches
   molecules are useful for diagnostic markers or therapeutic targets for the various diseases or morbid states. In particular, they are useful in gene therapy for treating osteoporosis, neurological disease, Alabaimer's disease, Parkinson's disease, dementia, short memory and various cancers, as well as for maintaining equilibrium of sense or motor function, and for treating emotional reaction, fear response and panic. Accordingly, they exhibit osteopathic, neuroprotective, nootropic, antiparkinsonian, cytostatic and tranquiliser activities. This polynucleotide is a full length human cDNA sequence of the invention. NOTE: This sequence is not given in the sequence listing of the specification but can be obtained on CD-ROM from the European Patent Office, Vienna Sub-office.
  This invention relates to novel, isolated full length human cDNA molecules and the encoded proteins thereof. Specifically, it refers to cDNA clones obtained by an oligo-capping method, where none of these clones are identical to any known human mRNAs. The present invention describes an immunoassay to identify agonists and antagonists, as well antibodies, antisense molecules and siRNAs that can all be used to bind antibodies, antisense molecules and siRNAs that can all be used to bind
  AAH18230;
  AAH18230 standard; cDNA;
  Sequence
   antibodies, antisense molecules and siRNAs that can all be to and modulate expression of the cDNA molecules. As such, molecules are useful for diagnostic markers or therapeutic
  Claim 1; SEQ ID
  New 1995 cDNA, useful for treating osteoporosis, neurological diseases, Alzheimer's diseases, Parkinson's diseases, dementia and various cancers.
  14-FEB-2003; 2003JP-00102207.
09-MAY-2003; 2003JP-00131452.
   18-AUG-2004
  EP1447413-A2
  Homo sapiens.
   osteopathic;
tranquiliser.
  Parkinson's disease; dementia; short memory; cancer;
sense or motor function; emotional reaction; fear response;
osteopathic; neuroprotective; nootropic; antiparkinsonian;
   Wakamatsu A,
  12-FEB-2004; 2004EP-00003145
  osteoporosis; neurological
   Full length human cDNA useful for treating neurological disease Seq 564.
  Isogai T,
   (REAS-) RES ASSOC BIOTECHNOLOGY
  2447
   3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  2004-583265/57.
  ss; human; oligo-capping method; diagnostic
   l Similarity
45; Conserv
   GTGCCACTGCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   2491 BP; 509 A; 639 C; 794 G; 549 T; 0 U; 0 Other;
   Yamamoto
A, Ishii
   Conservative
  NO 564; 2686pp; English.
  1.4%;
   ω Ć
   Nishikawa T, :
Nagai K, Irie
  3977
   disease; Alzheimer's disease;
   0;
  Score 45;
Pred. No.
   ВP
   Mismatches
   Isono Y, Sugiyama T,
e R;
  DB 13;
  .1e-10;
   <u>,</u>
  Length 2491;
   Indels
  marker;
  2491
  gene therapy;
   cytostatic;
   0
   Otsuki T;
  88
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Human cDNA sequence SEQ ID NO:18165

26-JUN-2001

(first entry)

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RESULT 140
AAK69446/c
ID AAK694
XX
AC AAK694
XX
DT 06-NOV
XX
DH Human
  닭
  S
   complementary strand of a polynucleotide which comprises a 5'-end sequence and an oligonucleotide comprises a 5'-end sequence and an oligonucleotide comprises a squence complementary to a polynucleotide which comprises a 5'-end sequence complementary to a polynucleotide which comprises a 3'-end sequence, where the oligonucleotide comprises at least 15 nucleotides and the combination of the 5'-end sequence/3'-end sequence is selected from those defined in the specification. The primer sets can be used in antisense therapy and in gene therapy. The primers are useful for synthesising polynucleotides, particularly full-length CDNAs. The primers are also useful for the detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs. The primers allow obtaining of the full-length CDNAs. The primers allow obtaining of the ANH13628 and CC cDNAs easily without any specialised methods. ANH03166 to ANH13628 and CC cDNAS and and and acid sequences; and ANH1363 to ANH13632 represent colligonucleotides, all of which are used in the exemplification of the
   Query Match
Best Local (
   Matches
   29-JUL-1999; 99JP-00248036.
27-AUG-1999; 99JP-00300253.
11-JAN-2000; 2000JP-00118776.
02-MAY-2000; 2000JP-00183767.
09-JUN-2000; 2000JP-00241899.
 Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24258
                         06-NOV-2001
   AAK69446;
  AAK69446
  Sequence 3977
  length cDNAs defined in the specification. Where a primer set comprises:
(a) an oligo-dT primer and an oligonucleotide complementary to the complementary strand of a polynucleotide which comprises one of the 5602 nucleotide sequences defined in the specification, where the oligonucleotide comprises at least 15 nucleotides; or (b) a combination
   Primer sets for synthesizing polynucleotides, particularly the 5602 full-
length cDNAs defined in the specification, and for the detection and/or
diagnosis of the abnormality of the proteins encoded by the full-length
  WPI; 2001-318749/34.
  28-JUL-2000; 2000EP-00116126
  07-FEB-2001.
  Human; primer;
  Claim 8;
  (HELI-) HELIX RES INST.
  Local Similarity
  3931
  present invention
   ω
  invention
  Isogai T,
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3975
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
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   Conservative
                        (first entry)
  BP; 1063 A; 888 C; 987 G; 1039 T; 0 U; 0 Other;
   detection; diagnosis; antisense therapy; gene therapy; ss
  Nishikawa T,
T, Wakamatsı
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   100.0%;
   1.4%; Score 45;
100.0%; Pred. No.
  describes primer sets for synthesising 5602 full-
   Wakamatsu A,
  4513
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  Hayashi K, S
A, Nagai K,
   Sequence Listing; English.
   DB 4; L
   Saito K,
K, Otsuki
   0
  Length 3977;
   Indels
  H
  Yamamoto
   0
   Gaps
   0
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```
Human; immune; haematopoietic; immune/haematopoietic antigen; cytostatic; gene therapy; vaccine; metastasis; ds.
   cancer;
```

WO200157182-A2

2001WO-US001354

2000US-0179065P

17-MAR-2000; 18-APR-2000; 19-MAY-2000; 28-JUN-2000; 02-MAR-2000; 16-MAR-2000; 2000US-0186628P 2000US-0186359P 2000US-0186359P 2000US-0186359P 2000US-0198123P 2000US-0214886P 2000US-0214886P 2000US-0216847P 2000US-021889P 2000US-021889P 2000US-021889P 2000US-021889P 2000US-021889P 2000US-021889P 2000US-0228519P 2000US-02252619P 2000US-02252619P 2000US-02252619P 2000US-02252619P 2000US-0225213P 2000US-0231447P 2000US-0231443P 2000US-0231443P 2000US-0231443P 2000US-0231443P 2000US-0231444P 2000US-0231443P 2000US-0231444P 2000US-0231444P 2000US-0231443P 2000US-0231443P 2000US-0231443P 2000US-0231443P 2000US-0231443P 2000US-0231443P 2000US-0231243P 2000US-0231243P 2000US-0231243P 2000US-0231243P 2000US-0231243P 2000US-0231243P 2000US-0231243P 2000US-0231243P 2000US-0231243P 2000US-0231249P 2000US-0231249P 2000US-0233063P 2000US-0233063P 2000US-0233064P

11-JUL-2000; 11-JUL-2000; 14-JUL-2000; 26-JUL-2000; 07-JUL-2000;

14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000; 14-AUG-2000;

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08-SEP-2000;

08-SEP-2000; 08-SEP-2000; 12-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000; 14-SEP-2000;

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RESULT 141
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XX AAL030
XX AAL030
DT 21-NOV
XX Human
XX Human
XX Homo s
XX Hom
  吊
   δ
  CC amino acid sequences given in AAMS2170 to AAMS1921. (I) have cytostatic cactivity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and crantent of diseases associated with inappropriate (I) expression. For ce example, they may be used to treat disorders associated with decreased ce expression by rectifying mutations or deletions in a patient's genome chat affect the activity of (I) by expressing inactive proteins or to concert affect the patients own production of (I). Additionally, (I) concert affect the patients own production of (I). Additionally, (I) concert acids into a host cell and culturing the cell to express the concert and treat immune/haematopoietic-related diseases, especially concerts and cancer metasses of haematopoietic acids express the concerts and cancer metasses of haematopoietic acids. AAK64703 cc cancers and cancer metasses of haematopoietic antigen genomic concerts antigen genomic sequences used in the exemplification of the present invention.
  Matches
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04-FEB-2000;
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02-MAR-2000;
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19-MAY-2000;
07-JUN-2000;
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05-JAN-2001; 2001US-0259678P.
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  02-AUG-2001.
  WO200155320-A2
  3073
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  2001-483426/52
  sapiens.
   ξ
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  for
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   acids encoding for preventing,
  gene therapy;
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCT
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2000US-0209467P.
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  Conservative
  2001WO-US001339
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  100.0%;
  24258; 3071pp + Sequence Listing; English
  1.4%;
   human immune/hematopoietic diagnosing and/or treating
  Ruben SM;
  INC
  ..
   ВP
  Score 45;
Pred. No.
   Mismatches
  DB 4;
3e-10;
  0
   T; 0 U; 0 Other;
   Length 4513;
   antigen polypeptides, cancers and metastasis.
   Indels
  2342
  3117
  5729.
   0
  AAM82169
invention
  Gaps
  and
  the
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2000US-0251198P.
2000US-0251198P.
2000US-0251185P.
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The present inventi
number of human rep
in the prevention a
including cancer. T
protein of the inve
   20-OCT-2000

20-OCT-2000

20-OCT-2000

20-OCT-2000

00-NOV-2000

01-NOV-2000

01-DEC-2000

  Isolated
used in F
Sequence 6565
  Disclosure;
  Rosen
   (HUMA-)
   preventing,
  HUMAN
   nucleic
  Barash
                              invention provides the protein and coding sequences cuman reproductive system related antigens. These can be ention and treatment of reproductive system disorders, ancer. The present sequence is a genomic sequence encothe invention
  SEQ
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2000US-0241809P.
2000US-0241809P.
2000US-0244617P.
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2000US-0249219P.
2000US-025919P.
2000US-02599P.
2000US-025919P.
2000US-02599P.
BP; 1349 A;
   GENOME
  Ħ
   acid molecul
ng, treating
  SC,
  No
  SCI
   reating or
  5729; 1297pp +
  Ruben
 1951
  SM;
   encoding a reproductive rameliorating a medical
Ç
 1910
  Sequence
<u>ი</u>
 1355
  Listing;
 Η,
 0
Ç;
```

0 Other;

encoding

be of a used a

English system and condition

antigen

18

28-JUN-2000
30-JUN-2000
31-JUN-2000
11-JUL-2000
11-JUL

20000US.

S-0214886P S-02174886P S-02176880P S-02279963P S-02229963P S-02252513P S-02252513P S-022525213P S-022525213P S-02252668P S-02252752P S-02252752P S-02252752P S-02252752P S-0225275P S-02252778P S-02252778P S-0225279P S-0225279P S-0225279P S-0235368P S-0235636P S-0235637P S-0235637P S-023637P S-0237939P S-023793P S-02379 S-023793P S-023793P S-023793P S-023793P S-023793P S-02379 S-023

| RESC<br>CXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | д <b>ж</b> в б                                                                                                                                                                                                                  |     |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              | Query Match Best Local Simila Matches 45; Co 9 3078 GTGCC 9 3078 GTGCC 9 5140 GTGCC                                                                                                                                             |     |
| DNA; 6565 BP.  t entry)  system related antigen DNA SEQ ID NG E System related antigen; reproductively; ds.  college system;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | 1.4%; Score 45; DB 4; Length 6565;  imilarity 100.0%; Pred. No. 3e-10;  Conservative 0; Mismatches 0; Indels 0; Gaps 0;  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 5096 |     |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |                                                                                                                                                                                                                                 |     |
| ***************************************                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      | מקל מקל מקל מקל<br>מאל מאל מאל מאל מאל מאל מאל מאל מאל מאל                                                                                                                                                                      | PR  |
| 14-SEP-2000<br>14-SEP-2000<br>14-SEP-2000<br>14-SEP-2000<br>14-SEP-2000<br>14-SEP-2000<br>21-SEP-2000<br>21-SEP-2000<br>21-SEP-2000<br>25-SEP-2000<br>25-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>29-SEP-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT-2000<br>20-OCT |                                                                                                                                                                                                                                 | 200 |
| 2000US-023239FP. 2000US-023239FP. 2000US-023239FP. 2000US-0232401P. 2000US-0232401P. 2000US-0232401P. 2000US-023499FP. 2000US-023499FP. 2000US-023499FP. 2000US-023499FP. 2000US-023499FP. 2000US-023499FP. 2000US-023548FP. 2000US-023536FP. 2000US-023636FP. 2000US-023636FP. 2000US-023636FP. 2000US-023636FP. 2000US-023636FP. 2000US-023636FP. 2000US-023636FP. 2000US-023636FP. 2000US-024637FP. 2000US-024178FP. 2000US-024178FP. 2000US-024677FP. 2000US-0246611P. 2000US-0249211P.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |                                                                                                                                                                                                                                 | 023 |

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RESULT 143
ABL97378/c
ID ABL97378;
XX ABL97378;
AC ABL97378;
XX ABL97378;
AC ABL97378;
AC ABL97378;
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XX Human; test
XX Human; test
XX Gardiovass
XX Gardiovass
XW Gardiovas
   밁
   δ
  Query Match
Best Local (
   Matches
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17-NOV-2000;
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17-NOV-2000;
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05-DEC-2000;
05-DEC-2000;
05-DEC-2000;
06-DEC-2000;
06-DEC-2000;
08-DEC-2000;
   The present invention provides the protein and coding sequences of a number of human reproductive system related antigens. These can be used in the prevention and treatment of reproductive system disorders, including cancer. The present sequence is a genomic sequence encoding a protein of the invention
                     17-JAN-2001; 2001WO-US001329
   02-AUG-2001.
   gastrointestinal
  reproductive system disorder; urinary system disorder; gene therapy; cardiovascular disorder; respiratory disorder; neurological disorder;
  Human; testicular antigen; testes; cancer; metastasis; immune disorder;
   Human testicular antigen encoding
   21-JUN-2002
  Sequence
  Disclosure; SEQ ID NO
  Isolated nucleic acid molecule encoding a reproductive system antigen used in preventing, treating or ameliorating a medical condition.
  Rosen
   WO200155317-A2
  (HUMA-)
   5140
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  δ
  HUMAN
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   BP; 1349 A; 1951 C; 1910
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  5730; 1297pp + Sequence Listing; English.
  Ruben
   6565
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%; Pred. No. 3e-
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   G; 1355
  DB 4;
3e-10;
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   ₫B.
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31-JAN-2000
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973 human testicular antigen polypeptides, useful sing and/or treating testicular cancer.
  RESULT 144
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XX 21-JUN
XX
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XX 1-JUN
XX
XX Human;
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XX
Y Cardio
XX
Y W02001
XX
Y 00-AUG
XX
Y 00-AUG
XX
Y 17-JAN
XX
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X 11-JUL
X R 11-JUL

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31-JAN-2000
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24-FEB-2000
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11-MAR-2000
17-MAR-2000
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07-JUN-2000
07-JUL-2000
07-JUL-2000
07-JUL-2000
11-JUL-2000
  The present invention provides the protein and coding sequences of 973 human testicular antigens, and fragments of their genomic sequences. The sequences can be used in the treatment of cardiovascular, urinary system, reproductive system, immune, respiratory, neurological and gastrointestinal disorders, infections, and particularly cancer, especially testicular cancers. The present sequence is a DNA encoding a
  Human; testicular antigen; testes; cancer; metastasis; immune disorder; reproductive system disorder; urinary system disorder; gene therapy; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disease; infection; cytostatic; gene; ds.
   Human testicular antigen encoding
   21-JUN-2002
  ABL97377;
   ABL97377
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   Disclosure;
  17-JAN-2001; 2001WO-US001329
  02-AUG-2001
  WO200155317-A2
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  1.4%; Scilarity 100.0%; F
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  DNA;
  6565
  Score 45; DB;
; Pred. No. 3e-
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  DB 4;
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  Length 6565
   Indels
   ID NO:
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   Gaps
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Nucleic acids encoding for preventing, diagnos WPI; 2001-483232/52.

HUMAN Barash

sc,

S

2001US-0259678P GENOME SCI INC

preventing,

diagnosing

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XX
AC AAX235
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08-NOV-2000
17-NOV-2000
17-NOV
   gastrointestinal disorders, infections, and especially testicular cancers. The present
  The present invention provides the protein and coding sequences of 973 human testicular antigens, and fragments of their genomic sequences. The sequences can be used in the treatment of cardiovascular, urinary system, reproductive system, immune, respiratory, neurological and
                  17-JUN-1999
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  Nucleic acids encoding 973 human testicular antigen polypeptides, useful for preventing, diagnosing and/or treating testicular cancer.
   Disclosure;
  (HUMA-) HUMAN GENOME
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s; Pred. No. 3e-
0; Mismatches
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  BP
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o. 3e-10;
o;
  Ģ.
  1355
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  U; 0 Other;
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   Gaps
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RESULT 146
AAS36670
  222222222222
UXAXEXBXBXBXBXBXB
  片
   S
  Query Match
Best Local
  atherosclerosis. The structure of AMF 18 used to traces substrates, e.g. for use in AmP assays. AmP, which hydrolyzes N-terminal imido bonds, can be used to degrade industrial protein feeds to free amino acids, to degrade proteinaceous wastes, as additives in enzyme formulations used to treat malabsorption syndrome and for studying its hinlogical role. Antibodies against AmP are used in immunohistochemical
  This invention describes the isolation of a novel human aminopeptidase P (Amp). This protein is used to produce recombinant Amp and can be used for gene therapy for treating Amp-deficiency conditions. Its fragments are used as primers and probes to identify patients with homozygous and heterozygous Amp deficiency, including prenatal diagnosis (patients defective in Amp are at risk of developing angioedema if treated with angiotensin-converting enzyme inhibitors), also as antisense inhibitors in cases of excessive Amp expression. The product of the invention is also used to identify Amp-expressing sequences in other animals and to generate transgenic animals, and comparisons of genomic sequences are used to detect mutations. Amp inhibitors are potentially useful as antihypertensive agents and to prevent or treat arterial (re)stenosis or atherosclerosis. The structure of Amp is used to design synthetic
               chicken; sheep; immunosuppressive; antiarthritic; vasotropic; dog; antirheumatic; antiproliferative; cytostatic; cardiant; neuroprotective; cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer;
   02-SEP-1997;
  02-SEP-1998;
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  arterial
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   Sequence 16595
   11-MAR-1999.
   Human kidney aminopeptidase P genimic DNA fragment
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  17-DEC-2001
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   Claim 13; Page 192-201; 201pp; English.
  (MEDI-)
  4301
   3078
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  MEDICAL COLLEGE GEORGIA RES
   Similarity
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   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
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  Conservative
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  BP; 4429 A; 4145 C;
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  98WO-US018426
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  100.0%;
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   B
  G; 3853
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   <u>ب</u>
  ٣
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  T, 0 U; 0 Other;
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  σ
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11-JUL-2000
11-JUC-2000
11-JUC
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hyperproliferative disorder; breast; liver; cardiovascular disorder; ds cerebrovascular disorder; nervous system disorder; bacterial infection; fungal infection; viral infection; occlar disorder; endocrine disorder; gastrointestinal disorder; respiratory disorder; wound healing; skin aging; organ transplantation; tissue regeneration; anti-infertility.

02-AUG-2001

17-JAN-2001; 2001WO-US001340

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S-0246476P.
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```
CC Cardiovascular system antigens and their associated polynucleotides are conservation in the diagnosis, treatment and prevention of various types of disorders in e.g. humans, mice, rabbits, goats, horses, cate, dogs, chickens or sheep. A pathological condition can be determined by contickens or sheep. A pathological condition can be determined by contickens or sheep. A pathological condition can be determined by contickens or sheep. A pathological condition can be determined by contickens or sheep. A pathological condition can be determined by contickens or sheep. A pathological condition can be determined by contickens or sheep. A pathological condition in a cardiovascular such as preservative disorders such as cardiovascular disorders such as neoplasms of the breast or liver, cardiovascular disorders such as cardiac arrest, cerebrovascular disorders such as cerebral ischaemia, concervations system disorders such as Alzheimer's disease, infections caused by bacteria, viruses and fungi, ocular disorders such as corneal confection, endocrine disorders such as premature labour and infertility, confection, endocrine disorders such as premature labour and infertility, confection, endocrine disorders such as premature labour and infertility, confection, endocrine such as Crohn's disease, renal disorders such as glomerulonephritis and respiratory disorders such as asthma and confection, to regenerate tissues and in chemotaxis. Note: The sequence data for this patent did not form part of the printed carter with out of the patent did not form part of the printed carter with out of the patent did not form at directly from WIPO as the printed carter such as obtained in electronic format directly from WIPO.
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08-DEC-2000;
08-DEC-2000;
11-DEC-2000;
05-JAN-2001;
  New cardiovascular system related polynucleotides and polypeptides, useful for diagnosing, treating and/or preventing disorders of the
   Sequences AAS35741-AAS36942 represent genomic
   Claim 1;
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  WPI; 2001-451930/48.
  (HUMA-) HUMAN GENOME SCI INC.
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2000US-0251990P.
2000US-0254097P.
2000US-0259678P.
   2000US-0251869P
  Ruben
   SM.
  DNA molecules, whi
  which
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encode

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닭
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  Query Match
Best Local :
   Matches
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4841
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100.0%; Pr
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AAK83984 standard; DNA; 17581 BP

07-NOV-2001 (first entry)

Human immune/haematopoietic antigen genomic sequence SEQ ID NO:38796.

Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis; ds.

## Homo sapiens.

WO200157182-A2

## 

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useful f
Disclosure; SEQ ID NO 38796; 3071pp + Sequence Listing; English.
  (HUMA-)
   2001-483426/52.
   Ş
                                  cacids encoding for preventing,
  HUMAN
   Barash
  2000US.
2000US.
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-025888888

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-02588
  SCI
                                  human immune/hematopoietic diagnosing and/or treating
   Ruben
  INC.
   X.
                                      antigen polypeptides, cancers and metastasis.
```

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X8888888888888888888
   밁
   S
   RESULT 148
   CC AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) camino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic CC activity, and can be used in gene therapy and vaccine production. (I) CC proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat discorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC polynucleotides may be used to produce the secreted (I), by inserting the CC nucleic acids into a host cell and culturing the cell to express the CC diagnose and treat immune/haematopoietic-derived cells. AAK64703 CC cancers and cancer metastases of haematopoietic antigen genomic CC sequences from the present invention. AAK54942 to AAK87694 represent invention. AAK54942 to AAK8769 and AAM82169
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07-JUN-2000;
30-JUN-2000;
07-JUL-2000;
07-JUL-2000;
11-JUL-2000;
11-JUL-2000;
11-JUL-2000;
  Human; cardiovascular system related polypeptide; cancer; proliferative disorder; foetal abnormality; developmental abnormality; haematopoietic disorder; ADES; autoimmune disease; rheumatoid arthritis; inflammation; allergy; neurological disorder; Alzheimer's disease; Parkinson's disease; cognitive disorder; schizophrenia; asthma; skin disorder; psoriasis; sepsis; diabetes; atherosclerosis; cardiovascular disorder; angiogenic disorder; kidney disorder;
   07-MAR-2002;
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  The invention relates to human cardiovascular system related polypeptides and the polynucleotides encoding them. The polypeptides, polynucleotides and antibodies to the polypeptides are useful for diagnosing a pathological condition or a susceptibility to a pathological condition, for preventing, treating, or ameliorating a medical condition, such as cancer of cardiovascular system tissues, proliferative disorders, foetal and developmental abnormalities, haematopoietic disorders, diseases of the immune system, AIDS, autroimmune diseases (e.g., rheumatoid the immune system, AIDS, autroimmune diseases (e.g., rheumatoid
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  Barash
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disorders

(e.g.,

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   diabetes, atherosclerosis, cardiovascular disorders, angiogenic disorders, kidney disorders, gastrointestinal disorders, pregnancy-related disorders, emocrine disorders and infections. The nuclesic acids are also useful for chromosome identification, radiation hybrid mapping or long-range restriction mapping. The polypeptides and polynuclectides may also be used as food additives or preservatives to increase or decrease storage capabilities, fat content or other nutritional components. This sequence represents human cardiovascular system related
   breast neoplasms; liver neoplasm; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; cerebrat ischaemia; angiogenesis; nervous system disorder; Alzheimer's disease; i ocular disorder; corneal infection; wound healing; epithelial cell proliferation; skin aging; sunburn;
   26-AUG-2002;
   08-JAN-2004
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   Gaps
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The invention relates to an isolated nucleic acid molecule encoding a CC human cardiovascular system associated polypeptide (or antigens), or its C fragment. Also included recombinant vectors, recombinant host cells, an CC isolated human cardiovascular system associated polypeptide (including CC its fragment, allelic variant, species homologue or epitope), an isolated CC antibody that binds specifically to a human cardiovascular system CC susceptibility to a pathological condition or CC susceptibility to a pathological condition comprising determining the CC susceptibility to a pathological condition based on the presence or absence of a mutation in human cardiovascular system CC associated nucleic acid and diagnosing a condition based on the presence CC or absence of the mutation), identifying a binding partner to human CC cardiovascular system associated polypeptides, the gene corresponding to the human cardiovascular system associated cDNA sequence and identifying an activity in a biological assay comprising expressing the human cardiovascular system associated cDNA in a cell, isolating the cardiovascular system associated cDNA in a cell, isolating the cardiovascular system associated cDNA in a cell, isolating the cardiovascular system associated cDNA in a cell, isolating the cardiovascular system associated cDNA in a cell, isolating the cardiovascular system associated nucleic acids and polypeptides are used the provent, treat or ameliorate a medical condition (for example in
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  2004-081713/08
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useful

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   SCI
   Ruben
  MS
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cardiac arrest, cerebrovascular disorders such as cerebral ischaemia, nervous system disorders such as Alzheimer's disease, infections caused by bacteria, viruses and fungi, ocular disorders such as corneal infection, endocrine disorders such as premature labour and infertility, gastrointestinal disorders such as gremature labour and infertility, gastrointestinal disorders such as Crohn's disease, renal disorders such as glomerulonephritis and respiratory disorders such as asthma and pleurisy. The polypeptides can also be used to aid wound healing, to prevent skin aging due to sunburn, to maintain organs before transplantation, to regenerate tissues and in chemotaxis. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published_pct_sequences
   Sequences AAS35741-AAS36942 represent genomic DNA molecules, which encode the cardiovascular system antigen polypeptides of the invention. Cardiovascular system antigens and their associated polynucleotides are useful in the diagnosis, treatment and prevention of various types of disorders in e.g. humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A pathological condition can be determined by detecting the presence or absence of a mutation in a cardiovascular system antigen polynucleotide. The treatable disorders include autoimmune diseases such as rheumatoid arthritis, hyperproliferative disorders such as reoplasms of the breast or liver, cardiovascular disorders such as cardiovascular system.
  New cardiovascular system related polynucleotides and polypeptides, useful for diagnosing, treating and/or preventing disorders of the
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  18-APR-2000;
19-MAY-2000;
07-JUN-2000;
28-JUN-2000;
  02-MAR-2000;
16-MAR-2000;
17-MAR-2000;
  Human; nootropic; neuroprotective; cytostatic; dermatological; viru immunosuppressive; antiinflammatory; anti-HIV; antibacterial; vulne antiparkinsonian; antisickling; antianaemic; antiarthritic; cancer; antirheumatic; hepatotropic; cerebroprotective; antiinflammatory; antiallergic; antidiabetic; antiulcer; anticonvulsant; antifungal; antiparasitic; cardiant; immune disorder; cardiovascular disorder;
  17-JAN-2001;
  WO200159063-A2
  Human nervous system related polynucleotide
   23-JAN-2002
  ABA15608;
  ABA15608
   16-AUG-2001.
   neurological disease;
   7806
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2000US-024921P

2000US-024921P

2000US-024921P
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
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  (HUMA-) HUMAN
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2000US-0251999P
2000US-0251999P
  GENOME
  sc,
   SCI
   Ruben
   83
```

2001-541565/60.

Nucleic acids encoding useful for preventing, metastases. 3224 human nervous system antigen polypeptides, diagnosing and/or treating nervous system cancers

Disclosure; SEQ ID ð 7939; 1701pp + Sequence Listing; English.

The invention relates to novel genes (ABA11004-ABA21534) and proteins CC (ABB14678-ABB18001) useful for preventing, treating or ameliorating CC medical conditions e.g. by protein or gene therapy. The genes are CC isolated from a range of human tissues disclosed in the specification. CC The nucleic acids, proteins, antibodies and (ant)agonists are useful in CC the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and CC ovarian cancer and other cancers of the adrenal gland, bone, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune CC disorders e.g. Addison's disease, allergies, autoimmune haemolytic CC annemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, CC multiple sclerosis, rheumatoid arthritis and ulcerative colitis; (c) CC cardiovascular disorders such as myocardial ischaemias; (d) wound healing (e) neurological diseases e.g. cerebral anoxia and epilepsy; and (f) CC infectious diseases such as viral, bacterial, fungal and parastic CC the printed specification, but was obtained in electronic format directly compared to the printed specification, but was obtained in electronic format directly WIPO at ftp.wipo.int/pub/published\_pct\_sequences

Sequence 17946 BP; 4718 A; 3994 C; 4320 G; 4914 T; 0 U; 0 Other;

밁 δ Query Match Best Local S Matches 45 7806 2898 GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 2942 ch 1.4%; Solid Similarity 100.0%; 1 45; Conservative 0; GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG Score 45; DB; Pred. No. 2.9 DB 5; 2.9e-10; Length 17946; Indels 0, Gaps 0

RESULT 152
ADB47506
ID ADB475
XX
AC ADB475
XX
AC ADB475
XX
AC 29-JAN
XX 29-JAN-2004 ADE47506; ADE47506 standard; (first entry) DNA; 17946 ВP

| B X X X X X X X X X X X X X X X X X X X                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |
|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Human; cardiovascular Human; cardiovascular Phuman; cardiovascular proliferative disord haematopoietic disord haematopoietic disord haematopoietic disord inflammation; allerg parkinson's disease; skin disorder; poori cardiovascular disorder; poori cardiovascular disorder; parkinson's disease; skin disorder; poori cardiovascular disorder; poori cardiovascular disorder; poori cardiovascular disorder; local parkinson's di                                                                                                                                                                                                                                                              |
| Human cardiovascular system related polypetid; Cancer, proditionative date date of the polypetid; Cancer, proditionative date of the polypetid; Cancer, proditionative date of the polypetid; Cancer, productive disorder; chicapteria, asthma; cardiovascular dasorder; papis; datasete; athrecoclerosis; personate, and polypetid; Cancer, productive disorder; kingy dasorder; personate, thingy dasorder; personate,                                                                                                                                                                                                                                                              |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |
| 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |
| 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 14-SEP-2000 21-SEP-2000 21-SEP-2000 25-SEP-2000 25-SEP-2000 26-SEP-2000 27-SEP-2000 29-SEP-2000 29-SEP-2000 29-SEP-2000 20-CCT-2000 21-NCV-2000 21-NCV                                                                                                                                                                                                                                                              |
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AC ADJ089
XX
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AC ADJ089
XX
DT 04-NOV
DT 04-NOV
DT Human
XX
DT Human
XX
English
XW
Dreast
XW
Dreast
XW
Cardia
  cc and the polymucleotides encoding them. The polypeptides, polymucleotides can antibodies to the polypeptides are useful for diagnosing a condition or a susceptibility to a pathological condition, can can entered for preventing, treating, or ameliorating a medical condition, such as cc cancer of cardiovascular system tissues, proliferative disorders, foetal can developmental abnormalities, haematopoietic disorders, diseases of ct immune system, AIDS, autoimmune diseases (e.g., rheumatoid arrhritis), inflammation, allergies, neurological disorders (e.g., carhritis), inflammation, allergies, neurological disorders (e.g., confict of disorders, asthma, skin disorders (e.g., psoriasis), sepsis, cc disorders, atherosclerosis, cardiovascular disorders, angiogenic cd disorders, atherosclerosis, cardiovascular disorders, pregnancy-cc disorders, atherosclerosis, cardiovascular disorders, pregnancy-cc are also useful for chromosome identification, radiation hybrid mapping correspondents. The nucleic acide care also useful for chromosome identification, radiation hybrid mapping correspondents. The nucleic acide care also the care capabilities, fat content or other nutritional components. This sequence represents human cardiovascular system related components. This sequence represents human cardiovascular system related
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08-DEC-2000;

11-DEC-2000;

11-DEC-2001;
   New cardiovascular system related polynucleotides and polypeptides, useful for preventing, treating, or ameliorating a medical condition, such as cancer of cardiovascular tissues and cancer metastases.
                   angiogenesis; nervous system disorder; Alzheimer's disease; ocular disorder; corneal infection; wound healing; epithelial cell proliferation; skin aging; sunburn;
   autoimmune disease; rheumatoid arthritis; hyperproliferative disorder; breast neoplasms; liver neoplasm; cardiovascular disorder; cardiac arrest; cerebrovascular disorder; cerebral ischaemia;
  Sequence 17946 BP; 4718 A;
   Human cardiovascular system associated polypeptide-related DNA SeqID2312
  04-NOV-2004
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  SEQ ID NO 2312; 262pp; English.
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2001US-0259678P.
2001US-00764869.
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  SM,
  1.4%;
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  human
  3994 C;
  0,
  Score 45;
Pred. No.
  cardiovascular system related
  Mismatches
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tissue regeneration; chemotaxis;
   B
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   10;
  4914 T;
  ç
   Length 17946;
  Indels
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  0 Other;
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11-JUL
  food additive; preservative; cardiovascular system associated antigen;
nuclear factor kappaB; NFkappaB; promoter element; human; ds.
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2000US-0251869P.
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2000US-0251990P.
   2000US-0234997P.
ACN45138
ID ACN4511
XX ACN4511
XX ACN451
XX ACN51
XX ACN5
   片
   δ
   The invention relates to an isolated nucleic acid molecule encoding a CC human cardiovascular system associated polypeptide (or antigens), or its CC fragment. Also included recombinant vectors, recombinant host cells, an CC isolated human cardiovascular system associated polypeptide (including its fragment, allelic variant, species homologue or epitope), an isolated CC antibody that binds specifically to a human cardiovascular system CC susceptibility to a pathological condition or CC susceptibility to a pathological condition based on the presence CC presence or absence of a mutation in human cardiovascular system CC associated pucleic acid and diagnosing a condition based on the presence CC or absence of the mutation), identifying a binding partner to human CC cardiovascular system associated polypeptides, the gene corresponding to CC the human cardiovascular system associated cDNA sequence and identifying CC an activity in a biological assay comprising expressing the human CC cardiovascular system associated cDNA in a cell, isolating the supernatant, detecting an activity in a biological assay and identifying CC supernatant, detecting an activity in a biological assay and identifying CC the protein in the supernatant having the activity. The human CC cardiovascular system associated mucleic acids and polypeptides are used to prevent, treat or ameliorate a medical condition (for example in CC humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep), for cxample autoimune diseases such as rheumacoid arthritis,

CC hyperproliferative disorders, for example neoplasms of the breast or liver, cardiovascular disorders, for example cardiac arrest,
   Query Match
Best Local Similarity
Matches 45; Conserv
  05-JAN-2001;
17-JAN-2001;
07-MAR-2002;
  01-MAR-2002; 2002US-00087192
   18-NOV-2004
   ACN45138;
  ACN45138 standard; DNA; 23694 BP
   28-FEB-2003; 2003WO-US006235.
   12-SEP-2003
  WO2003073826-A2
  Homo sapiens.
   Cytostatic; carcinoma; lymphoma; cancer; human;
   Human genomic sequence hCG17175.
   New cardiovascular system-related nucleic acid molecule, useful for diagnosing, preventing or treating diseases of the cardiovascular system, and in chromosome mapping, drug screening or in pharmacogenomics.
                                   (SAGR-) SAGRES DISCOVERY
   Disclosure; SEQ ID NO 2312; 262pp; English.
   (HUMA-) HUMAN GENOME SCI INC
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  7806 GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 7850
  2004-081713/08
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   Ruben
   1.4%; So llarity 100.0%; F Conservative 0;
  2001US-0259678P.
2001US-00764869.
2002US-00091504.
   (first entry)
   MS,
  Barash
   Score 45;
Pred. No.
   SC
  Mismatches
  DB 13,
3. 2.9e-10;
0;
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Length 17946; Indels

0

Gaps

0

gene;

88

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ACN44954
XX
AC ACN4
AC ACN4
XX
DT 18-1
XX
Cyt
XX
Hom
XX
HO
   S
  RESULT 155
   밁
   Query Match
   are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bloactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of bloactive agent capable of modulating the activity of CAP; (iv) for a bloactive agent capable of modulating the activity of CAP; (iv) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing carcinoma; (vii) for inhibiting the activity of CAP; (vii) for treating carcinoma; (viii) for neutralizing the effect of CAP; (ix) as a blochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent US2002182586A1, for which no sequence data was published
The present invention relates to novel DNA and protein sequences which are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bloactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of a bioactive agent capable of modulating the activity of CAP; (iv) for
  Claim 1; SEQ ID NO 1936; Opp; English
   Human genomic sequence hCG38622.
   ACN44954;
  The present invention relates to novel DNA and protein sequences
  comprises a nucleotide sequence.
   Recombinant nucleic acid useful for diagnosis and treatment of carcinoma
  Recombinant nucleic acid useful for diagnosis and treatment of carcinoma
  WPI; 2003-328604/31.
  Morris
   01-MAR-2002; 2002US-00087192
  28-FEB-2003; 2003WO-US006235
  12-SEP-2003.
  WO2003073826-A2
   Homo sapiens
  Cytostatic; carcinoma; lymphoma; cancer; human; gene; ss.
  18-NOV-2004
  ACN44954 standard; DNA; 31116 BP
   Sequence 23694 BP; 5742 A; 6329 C; 5736 G; 5781 T;
   Claim 1; SEQ
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  Local
  18194
  2003-328604/31
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  ¥
  Similarity
  TGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT 18238
   TGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT 3121
   nucleotide
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   (first entry)
  DISCOVERY
   100.0%; Pred. No. 2.9
cive 0; Mismatches
  1.4%;
  sequence
  Opp; English.
  Score 45;
Pred. No.
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   Length 23694;
   Indels
  0 U; 106 Other;
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   Gaps
  which
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RESULT 156
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ID ADZ132
XX ADZ132
XX Diagno
XW Cytost
OS Homo s
XX Diagno
XW Cytost
XX Diagno
XW Cytost
XX PO 07-APR
XX WO2005
XX WO2005
XX WO16
PF 23-SEF
XX WO18
PF 23-SEF
XX WO11;
ZY MOTIS
ZX MOTIS
ZX WPI; 2
ZX Nucle
PT compri
XX Diagno
XX Diacl
XX WPI; 2
ZX Nucle
PT compri
XX Diacl
XX Diacl
XX WIS Diacl
XX WIS Diacl
XX Diacl

  នននិននិនិនិង
   片
   5
                           CC associated (CA) nucleic acid, comprising two or more nucleic acid probes.

CC rhe invention also relates to a peptide array comprising two or more solution also relates to a peptide array comprising two or more composed by a CA nucleic acid sequence, a compound composed that binds to a polypeptide, an isolated antibody or its fragment which composition comprising the polypeptide or its antigen binding composition comprising the polypeptide or its antigen binding composition comprising the polypeptide or its antigen binding composition comprising the composition comprising the antibody and a carrier, a method of fragment, a composition comprising the composition of accountibody and a carrier, a method of screening for anticancer activity, a composition of accounting a CA nucleic acid, a method of diagnosing cancer, a composition of a CA nucleic acid in a cell. The CA nucleic acids are useful for detecting CA composition acids. The antibody is useful for detecting the presence or cancer cells in an individual which involves contacting cells from the individual with the antibody and detecting a complex of a CA composition from the cancer cells and the antibody, where the detection of the complex contacting cells are useful from the cancer cells and the antibody, where the detection of the complex contacting cells are useful for activity and the antibody, where the detection of the complex contacting cells are useful to the presence of cancer cells and the antibody, where the detection of the complex contacting cells are useful to the presence of cancer cells and the antibody.
   Matches
   Best
  Query Match
  evaluating the effect of a candidate carcinoma drug; (v) for diagnosing carcinoma; (vi) for inhibiting the activity of CAP; (vi) for treating carcinoma; (vii) for neutralizing the effect of CAP; (vi) as a blochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent US2002182586A1, for which no sequence data was published
  Diagnosis; DNA microarray; microarray; biochip; cancer; neoplasm;
  Human cancer-associated genomic DNA #63
  16-JUN-2005
   ADZ13255;
   ADZ13255 standard;
  Sequence 31116 BP; 7214 A; 8217 C; 7722 G; 7963 T; 0 U; 0 Other;
   Disclosure; SEQ ID NO 775; 198pp; English
   comprises two
  Nucleic acid array useful for detecting cancer associated nucleic acid,
   WPI; 2005-273395/28
   Morris
   23-SEP-2003; 2003US-00669920
  23-SEP-2004; 2004WO-US031617
  WO2005031001-A2
  Homo sapiens
  cytostatic; gene; ds.
  The invention relates to a nucleic acid array for detecting a cancer
  Local
  3078 GTGCCACTGCACTCCAGCCTGGGCAACACAGACAAGACTCTGTCTC 3122
   6981 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   l Similarity
45; Conserv
   DW, Malandro
               correlates with the
   Conservative
  (first entry)
   or more nucleic acid probes
   DNA;
   MS
  100.0%;
  1.48;
   31279
   ٥,
  Score 45;
Pred. No.
   ₽P
   Mismatches
               presence of cancer
   2.8e-10;
  DB 11;
  Length 31116;
   Indels
   0,
   Gaps
```

individual.

The composition is useful for inhibiting growth of cancer

```
ន្តន្តន្តន្តន្ត្
S
  밁
  S
  Query Match
Best Local S
Matches 45
                               Matches
  Query Match
Best Local Similarity
  The present invention relates to novel DNA and protein sequences which are associated with carcinomas. The sequences are useful for: (i) for screening drug candidates; (ii) for screening of bioactive agent capable of binding to Carcinoma Associated Protein (CAP); (iii) for screening of a bioactive agent capable of modulating the activity of CAP; (iv) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing carcinoma; (vii) for inhibiting the activity of CAP; (ix) as a biochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent respects to the carcinoma and the CAP are useful as markers of carcinoma capacity.
  Recombinant nucleic acid useful for diagnosis and treatment of carcinoma comprises a nucleotide sequence.
   Human genomic sequence
   ACN45014 standard;
  Sequence 31279 BP; 7246 A; 8268 C;
  cells in an individual or for delivering a therapeutic agent to cancer cells in an individual. The invention is also useful for diagnosing cancer, for treating cancer and for inhibiting expression of a CA gene in a cell. This sequence represents human cancer-associated genomic DNA of
  Sequence 32706
   01-MAR-2002; 2002US-00087192
   28-FEB-2003; 2003WO-US006235
   12-SEP-2003
   Homo sapiens
   Cytostatic; carcinoma; lymphoma; cancer; human; gene; ss.
   18-NOV-2004
   ACN45014;
  US2002182586A1,
  Claim 1; SEQ ID NO 1750; Opp; English.
   WO2003073826-A2
  (SAGR-)
  7137
2898 GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 2942
  3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  2003-328604/31
                               45;
  SAGRES
   Similarity
   GTGCCACTGCAGCTGCGGCAACAGAGCAAGACTCTGTCTC
                               Conservative
  Conservative
   (first entry)
  BP; 8225 A;
  DISCOVERY
  for which no sequence data was published
   DNA;
   1.4%;
   100.0%;
   1.4%; Score 45; DB 14;
100.0%; Pred. No. 2.8e-10;
   hCG14907
   32706
  7861 C; 8277 G; 8343 T; 0 U; 0 Other;
                              0,
  <u>.</u>
   Score 45;
Pred. No.
                               Mismatches
  Mismatches
  7755
   DB 11; I
2.8e-10;
   G; 8010 T; 0 U; 0 Other;
                              0
  0
   Length 31279;
  Length 32706;
                               Indels
  Indels
  7181
                              0;
  ٥,
                              Gaps
  Gaps
                               0
  0
```

```
RESULT 158
ADL82795/c
SXEXXXXXXXX
  밁
   S
   밁
  1.4%;
Best Local Similarity 100.0%;
Matches 45; Conservation
  The invention relates to a method of inhibiting the proliferation of a cancer cell comprises contacting the cell with a semaphorin3B (SEMA3B) polypeptide. The composition and methods are useful in diagnosing or treating cancer. The SEMA3B polypeptide inhibits tumour growth and induces apoptosis in cancer cells. The present sequence represents DNA encoding human semaphorin3B, SEMA3B.
                      SNP detection; diagnosis; non-insulin dependent diabetes; antidiabetic; anorectic; endocrine disease; gastrointestimetabolic disorder; nutritional disorder; gene; ds.
   08-SEP-2005
   Sequence 36534 BP; 7493 A; 10597 C; 10438 G; 8006 T; 0 U; 0 Other;
   Disclosure; SEQ ID NO 3; 75pp; English
  Inhibiting the proliferation of a cancer cell (e.g. breast cancer cell, lung cancer cell or prostate cancer cell) comprises contacting the cell with a semaphorin3B polypeptide that suppresses tumor growth.
  31-OCT-2001; 2001US-0335783P
   31-OCT-2002; 2002US-00285351
   cancer cell proliferation; semaphorin3B; SEMA3B; cancer; tumour growth;
  Human semaphorin3B,
  Human genomic
  AEB32373
  P-PSDB;
   Minna J,
  04-SEP-2003.
   US2003166557-A1
  Homo sapiens
   apoptosis; human;
   20-MAY-2004
  ADL82795;
   ADL82795 standard;
  (TEXA)
  15591
  7478
  3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   2003-898098/82
)B; ADL82793.
  UNIV TEXAS SYSTEM
   GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
  standard; DNA; 38678
   Tomizawa Y,
  (first entry
   (first
  DNA #14.
  ds; gene
   DNA; 36534
  SEMA3B,
   entry)
   Sekido
  DNA.
   0;
  Score 45; DB 11; 1
Pred. No. 2.8e-10;
   ВP
   ₽₽
   ۲
   Mismatches
   Lerman
                                      gastrointestinal disease;
  Length 36534;
  Indels
  15635
   0
   Gaps
```

0

Homo sapiens

```
ARBSULT 160
ARB32391/c
ID ARB323
XX
AC ARB323
XX
DT 08-SEP
XX
DE Human
XX
XX
DX SNP de
  S
   밁
  CC its sub-region, a computer readable medium having stored in it the SNP CC relational information given in the specification, an isolated nucleic acid molecule for detecting at least one SNP given in the specification CC comprising at least about 12 contiguous nucleotides, genotyping at least one SNP position given in the specification in the specification comprising at least one container containing a disorder and a kit CC comprising at least one container containing the detection reagent. CC Determining whether a trait is linked to one of the human chromosomes or its sub-region comprises determining whether the trait is linked to one cC or more SNPs using the detection reagents. Genotyping at least one SNP position given in the specification in a sample comprises contacting the callels at at least one SNP position given in the specification and cC allels at at least one SNP position given in the specification, and cC determining which allele is present at the at least one SNP position. CC comprises genotyping at least one SNP given in the specification in a conclusive genotyping at least one SNP given in the specification in a conclusive acid sample from the individual. The disorder is type II diabetes conclusive particularly type II diabetes or obesity. The detection reagent is cC useful in identifying an individual having or at risk of developing a cC represents human genomic DNA used in the scope of the invention. Note: CC represents human genomic DNA used in the scope of the printed cC specification but was obtained in electronic format from USPTO at tx
   Query Match
Best Local :
  Matches
  Venter JC, Zuca--
   The invention relates to a detection reagent capable of detecting one more single nucleic acid polymorphisms. The invention also relates to determining whether a trait is linked to one of the human chromosomes
   New detection reagent capable of detecting 1, 100, 500, 1000 or 5000 or more single nucleic acid polymorphisms, useful in identifying an
  08-SEP-2000; 2000US-0231397P.
10-SEP-2001; 2001US-00948947.
                                      Human genomic DNA #32.
  Disclosure; SEQ ID NO 136; 31pp; English
  WPI; 2005-511776/52.
  US2005147987-A1
SNP detection; diagnosis; non-insulin dependent diabetes; obesity;
   08-SEP-2005
   ABB32391;
  ABB32391 standard; DNA; 38684
   Sequence 38678 BP; 9340 A; 9040 C; 9074 G; 10537 T; 0 U; 687 Other;
   individual having or at risk of developing type II diabetes or obesity.
   (APPL-)
   19-JUL-2004; 2004US-00893315
   Local
   23579
  2895 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACA 2939
  45
   APPLERA CORP
   Similarity
   GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACA 23535
   Zhang JN,
  Conservative
   (first entry)
  ຸດ
   Z
   100.0%;
   1.4%;
   Liu
  Woodage
   X, Rowe W, Cravchik A,
  ..
   Score 45;
Pred. No.
  Mismatches
   DB 14;
   2.8e-10;
  0
   Length 38678;
  Indels
   Kalush
  0,
  Gaps
   one or
   유
  0
```

```
ADN31618
ID ADN3
XX
AC ADN3
XX
DT 12-1
  CC relational information given in the specification, an isolated nucleic comprising at least about 12 contiguous nucleotides, genotyping at least one SNP position given in the specification in a sample, identifying an comprising at least one container containing the detection reagent. CC comprising at least one container containing the detection reagent. CC comprising at least one container containing the detection reagent. CC its sub-region comprises determining whether the trait is linked to one or more SNPs using the detection reagents. Genotyping at least one containing whether the trait is linked to one cor more SNPs using the detection reagents. Genotyping at least one SNP contacting the containing which allele is present at the steeperification, and cc determining which allele is present at the at least one SNP position. CC comprises genotyping at least one SNP given in the specification in a contacting the comprises genotyping at least one SNP given in the specification in a contacting the comprises genotyping at least one SNP gosition. CC comprises genotyping at least one SNP given in the specification in a contacting the comprises genotyping at least one SNP given in the specification in a contacting the comprises genotyping at least one SNP given in the specification in a contacting the comprises genotyping at least one SNP given in the specification in a contact in strip in dispetes or obesity. The detection reagent is useful in identifying an individual having or at risk of developing a disorder particularly type II disabetes or obesity. This sequence contaction but was obtained in the scope of the printed contaction but was obtained in electronic format from USPTO at
   밁
   5
  RESULT 161
   Best
  Query Match
  New detection reagent capable of detecting 1, 100, 500, 1000 or 5000 or more single nucleic acid polymorphisms, useful in identifying an individual having or at risk of developing type II diabetes or obesity.
  Sequence 38684 BP;
  The invention relates to a detection reagent capable of detecting one or more single nucleic acid polymorphisms. The invention also relates to determining whether a trait is linked to one of the human chromosomes or its sub-region, a computer readable medium having stored in it the SNP
   Disclosure; SEQ ID NO 154; 31pp; English.
  08-SEP-2000; 2000US-0231397P.
10-SEP-2001; 2001US-00948947.
  07-JUL-2005
  antidiabetic; anorectic; endocrine disease; gastrointestinal disease; metabolic disorder; nutritional disorder; gene; ds.
   seqdata.uspto.gov/sequence.html.
   19-JUL-2004; 2004US-00893315
   US2005147987-A1
  (APPL-) APPLERA CORP NY
   Local Similarity
  23584 GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACA 23540
  2895
  JC, Zhang JN,
, Subramanian G,
  GGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACA 2939
   Conservative
  9340 A; 9042 C;
   100.0%;
  1.4%;
   Liu X, Ro
;, Woodage
   0;
   Score 45;
Pred. No.
   Rowe W,
lage T;
   Mismatches
   9075
   DB 14;
2.8e-10;
   G; 10540 T; 0 U; 687 Other;
   Cravchik A,
  14;
  Length 38684;
   Indels
   Kalush
   0,
   Gaps
```

12-AUG-2004 ADN31618;

(first entry)

ADN31618 standard; DNA; 39566 BP

```
RESULT 162
ABX14652
ID ABX146
XX ABX146
XC ABX146
XX Human
XX Human;
KW Human;
KW cardic
XX Cos Homo s
XX Homo s
XX Key
FH Key
FT Variat
   밁
   S
  Query Match
Best Local
  Matches
   The invention relates to a new compound 8-80 nucleobases in length (an antisense oligonucleotide) targeted to a nucleic acid molecule encoding squalene synthase (also known as farnesyl diphosphate farnesyl transferase 1), where the compound specifically hybridises with the nucleic acid molecule encoding human squalene synthase appearing as ADN31611 and inhibits the expression of squalene synthase in cells or tissues, screening for a modulator of squalene synthase, a diagnostic method for identifying a disease state, a kit or assay device comprising the compound and treating an animal having a disease or condition associated with squalene synthase. The compound and methods are useful in diagnosing and treating disorders related to cholesterol biosynthesis e.g. atheroselerosis, coronary heart disease and hypercholesterolaemia. The present sequence is a squalene synthase genomic DNA sequence, a target for the antisense oligonylatical
   Human gene encoding squalene synthase.
   05-MAR-2003
  Sequence
   New oligonucleotide targeted to a nucleic acid molecule encoding squalene synthase, useful in diagnosing and treating atherosclerosis.
   23-NOV-2002; 2002US-00304125
   23-NOV-2002; 2002US-00304125
   27-MAY-2004
   US2004102405-A1
   Homo
  Human; ds; antisense; farnesyl diphosphate atherosclerosis; corc
  Homo sapiens
  cardiovascular
   Human; ds;
   ABX14652;
  WPI; 2004-399735/37.
   Human squalene
            variation
   ABX14652 standard;
   (ISIS-) ISIS PHARM INC
  Local
  25338
   the antisense oligonucleotides.
  2898 GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 2942
  sapiens
  45;
  Similarity
  GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 25382
   39566 BP;
   antisense;
  gene; squalene synthase; cholesterol-related disease;
ılar disease; chromosome 8.
   Bennett CF,
   Conservative 0;
  SEQ ID NO
   (first
   synthase
 Location/Qualifiers replace(825,A) /*tag= o
   ense; squalene synthase;
hate farnesyl transferase 1; cholesterol;
coronary heart disease; hypercholesterolaemia.
   DNA;
   9928 A;
   entry)
  11; 67pp;
   40090
   genomic
   Dean
  8277 C;
  Score 45;
Pred. No.
   ₽P
  MX,
  DNA
  Mismatches
   English
   9254
  Dobie KW;
   DB 12; I
2.8e-10;
hes 0;
  <u>ი</u>
  12107
  Length 39566;
  Τ,
   Indels
  0 U; 0 Other;
   ;
   Gaps
   0
   exon
   variation
   variation
  variation
  variation
   variation
   variation
  variation
  variation
  variation
   variation
  exon
   SdC
   variation
  exon
   variation
  intron
  variation
   variation
  variation
  variation
  variation
  intron
                       intron
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9054. .25147
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replace(7398,C)
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replace(7355,T)
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   replace (4889, A)
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replace(4887,T)
   replace (4886, C)
  replace (4791, T)
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   2058. .2156
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   ceplace (8873,T)
   'number=
  number=
   standard_name=
   ceplace (5110, T)
   replace (2632, T)
  product= "Squalene
   standard_name=
2058. .37739
   870. .9053
   standard_
   095. .8869
  eplace (8031,G)
   standard_name=
  standard_name=
  standard_name=
  standard name=
   standard_name=
   =gar
   standard_name=
  997. .8094
   *tag=
   eplace (4430,C)
  standard name=
   *tag=
   æ
  쁑
  ac
   name=
  "Single nucleotide polymorphism"
                                 "Single nucleotide polymorphism"
   "Single nucleotide polymorphism"
  "Single nucleotide polymorphism"
  "Single
  "Single
  "Single
   "Single nucleotide polymorphism"
   "Single nucleotide polymorphism"
  "Single
   "Single
  "Single nucleotide polymorphism"
  "Single
  "Single
   "Single
  "Single
   "Single
   "Single
  synthase"
   nucleotide polymorphism"
  nucleotide polymorphism"
   nucleotide polymorphism"
  nucleotide polymorphism'
  nucleotide polymorphism"
   nucleotide polymorphism"
   nucleotide
  nucleotide
  nucleotide polymorphism<sup>n</sup>
  nucleotide polymorphism"
  nucleotide
  polymorphism"
   polymorphism"
  polymorphism"
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polymorphism"

polymorphism"

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variation
  variation
 exon
   exon
  variation
  variation
   exon
  variation
  variation
  variation
  variation
  variation
  variation
                        variation
   variation
  intron
   variation
   variation
   variation
  variation
  variation
  intron
   variation
  variation
  variation
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replace(22230,T)
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replace(20362,A)
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replace(27151,C)
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9366. .29542
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  standard_name= "Single
   standard_name=
eplace(26625,G)
  standard_name= "Single
eplace(26078,A)
  eplace (25686, A)
   number=
   standard_name=
  :ag= an
   :ag= am
ard name= "Single .30792
  ah
   25339
                               name= "Single
  "Single
  "Single
   "Single
  "Single
   "Single
   "Single
   "Single
  "Single
   "Single
  "Single
   "Single
  "Single
   "Single
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  polymorphism"
   polymorphism'
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   polymorphism
   polymorphism'
  polymorphism
   polymorphism"
   polymorphism"
   polymorphism'
   polymorphism'
   polymorphism'
  polymorphism
                                polymorphism"
   polymorphism'
   polymorphism'
  polymorphism'
RRSULT 163
ADN95863
ID ADN958
XX ADN968
XX ADN968
XX AOVel
DT 26-AUG
XX diseas
XW enzyme
XX Homo s
XX Homo s
XX FH Key
FH Key
FT Variat
FT CDS
FT CDS
FT EXON
  밁
   *************************************
   8
  Matches
   Query Match
Best Local Similarity
   disease diagnosis; gene eapression associated enzyme peptide; human; gene; ds.
  variation
   variation
   variation
   Homo sapiens
   Novel human
   26-AUG-2004
  ADN96863;
  ADN96863
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  variation
  variation
  variation
  variation
  variation
   variation
   variation
  intron
  variation
  25713
  2898
   45;
  GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
  standard;
  GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG
   enzyme
   Conservative 0;
   (first entry)
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replace(30732,C)
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replace(36607,G)
/*tag= bk
/standard_name= '
   /standard_name=
30793. .37517
/*tag= m
   /*tag=
   Location/Qualifiers replace (825, A)
  /standard_name=
replace(36541,C)
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replace(34532,C)
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replace(34451,C)
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replace(34249,T)
  /standard_name=
replace(32525,G)
  /standard_name=
replace(34179,T)
  replace (32032, A)
  standard_name=
   standard
  *tag= bc
standard_name=
  standard
   eplace (30841,G)
   eplace (31376,A)
   genomic
  DNA;
   40090
  name=
  DNA.
   Score 45; DB; Pred. No. 2.8
   ВP
   "Single
  "Single
   "Single
   "Single
  "Single
  "Single nucleotide
  "Single
  "Single
  "Single
  "Single
   "Single
   "Single
                          human
   DB 8; L-
5. 2.8e-10;
0;
   nucleotide polymorphism'
  nucleotide polymorphism"
   nucleotide polymorphism'
   nucleotide polymorphism"
  nucleotide polymorphism"
   nucleotide polymorphism
  nucleotide polymorphism"
  nucleotide polymorphism"
  nucleotide polymorphism"
  nucleotide
  nucleotide
  disorder;
```

Length 40090; Indels

2942 25757

0

Gaps

0

polymorphism"

gene

expression,

| FT variation             | FT variation                | FT variation                                                 | tron                                                       |                                                                     | FT exon                                               | variati                                                       |                                                            | FT intron<br>FT<br>FT                                                     | FT variation<br>FT<br>FT                                         | FT exon                             |                            | FT<br>FT variation                                               | FT<br>FT variation                                     | FT variation                                           | FT variation                       | FT<br>FT variation                                     | variat                                                 | variati                                                | at i                                                   | FT<br>FT variation                                     | FT<br>FT variation                                               | FT variation                                      | FT variation                           | FT intron |
|--------------------------|-----------------------------|--------------------------------------------------------------|------------------------------------------------------------|---------------------------------------------------------------------|-------------------------------------------------------|---------------------------------------------------------------|------------------------------------------------------------|---------------------------------------------------------------------------|------------------------------------------------------------------|-------------------------------------|----------------------------|------------------------------------------------------------------|--------------------------------------------------------|--------------------------------------------------------|------------------------------------|--------------------------------------------------------|--------------------------------------------------------|--------------------------------------------------------|--------------------------------------------------------|--------------------------------------------------------|------------------------------------------------------------------|---------------------------------------------------|----------------------------------------|-----------|
| ard name= "<br>(10460,C) | ard_name= "<br>e(9847,T)    | /:cay- y<br>/number= 3<br>replace(9190,T)<br>/*tag= 7        | x<br>rd_name= '<br>25147                                   | G H .                                                               | -                                                     | replace (8310,A) /*tag= u /standard_name= "S replace (8462 G) | "_                                                         | . 9                                                                       | 1,G)<br>ame= "                                                   | /9978094<br>/*tag= q<br>/number= 2  | tag= p<br>tandard_name= "  | <pre>/*tag= o /standard_name= "S replace(7653,C)</pre>           | <pre>/*tag= n /standard_name= "S replace(7398,C)</pre> | <pre>/*tag= m /standard_name= "S replace(7355,T)</pre> | e 2                                | <pre>/*tag= k /standard_name= "S replace(6911,A)</pre> | <pre>/*tag= j /standard_name= "S replace(5110,T)</pre> | <pre>/*tag= i /standard_name= "S replace(4889,A)</pre> | <pre>/*tag= h /standard name= "S replace(4887,T)</pre> | <pre>/*tag= g /standard_name= "S replace(4886,C)</pre> | <pre>/*tag= f /standard name= "S replace(4791,T)</pre>           | e<br>ard_name= "<br>e(4430,C)                     |                                        | 21577996  |
| Single nucleotide polymo | Single nucleotide polymo    |                                                              | 'Single nucleotide polymo                                  |                                                                     | Single nucleotide polymo                              | nucleotide                                                    | "Single nucleotide polymo                                  |                                                                           | Single nucleotide polymo                                         |                                     | Single nucleotide polymo   | ingle nucleotide                                                 | Single nucleotide polymo                               | ingle nucleotide                                       | ingle nucleotide                   | ingle nucleotide                                       | ingle nucleotide                                       | ingle nucleotide                                       | ingle nucleotide                                       | ingle nucleotide                                       | ingle nucleotide                                                 | Single nucleotide polymo                          |                                        |           |
| polymorphism"            | polymorphism"               |                                                              | polymorphism"                                              |                                                                     | polymorphism"                                         | polymorphism"                                                 | polymorphism"                                              |                                                                           | polymorphism"                                                    |                                     | polymorphism"              | polymorphism"                                                    | polymorphism"                                          | polymorphism"                                          | polymorphism"                      | polymorphism"                                          | polymorphism"                                          | polymorphism"                                          | polymorphism"                                          | polymorphism"                                          | polymorphism"                                                    | polymorphism"                                     |                                        |           |
| FT.                      | 7 H F                       | 자 당 한 1                                                      | 2 F F F T                                                  | Ld<br>Ld                                                            | 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7                 | 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7                         | FT                                                         | F F F                                                                     | FFT                                                              | P P P                               | FT                         | FT FT                                                            | P P P                                                  | F F F                                                  | 1 7 7 7<br>1 1 1 1                 | 77 T                                                   | FT<br>FT<br>FT                                         | 11.1<br>11.1<br>11.1                                   | P P P                                                  | P P P                                                  | P 7 7                                                            | P P P                                             | ************************************** | FT        |
| variation                | intron                      | variation                                                    | exon                                                       | variation                                                           | variation                                             | intron                                                        | -                                                          | variation                                                                 | variation                                                        | variation                           | variation                  | variation                                                        | variation                                              | variation                                              | intron                             | exon                                                   | variation                                              | variation                                              | variation                                              | variation                                              | variation                                                        | variation                                         | variation                              |           |
| replace(<br>/*tag=       | 30793<br>/*tag=<br>/number= | replace<br>/*tag=<br>/standa                                 | /standar                                                   | /stand                                                              | /number=<br>replace(2                                 | /*tag=<br>/number=<br>29543                                   | /*tag= a:<br>/standard<br>293662                           | /*tag= ac<br>/standard<br>replace(2)                                      | /*tag=<br>/standareplace                                         | /*tag=<br>/standare                 | /stai                      | /standa<br>replace<br>/*tag=                                     | /standa<br>replace<br>/*tag=                           | replace<br>/*tag=                                      | /nu<br>/*t                         | /st<br>251<br>/*t                                      | rel<br>/st                                             | rej<br>/st                                             | /et<br>/ep                                             | /*#<br>Kep /<br>## / ##                                | rep<br>/*t                                                       | ref<br>/*t                                        | rep<br>/*t                             | , *·      |
| 30841,G                  | .37517<br>ay<br>r= 6        | eplace(30732,C)<br>*tag= ax<br>standard name=                | rd_name=<br>.30792<br>aw                                   | ard name=<br>e(29761,T<br>av                                        | er= 5<br>ce(29572,T                                   | er= 5<br>30639                                                | = ar<br>dard_name=<br>29542                                | = aq<br>dard_name=<br>ce(28772,A                                          | *tag= ap<br> standard_name=<br> eplace(28032,A                   | ;= ao<br> dard_name=<br> ce(27151,C | ndard name=<br>ace(26625,G | standard_name=<br>eplace(26078,A<br>*taq= an                     | indard name=<br>lace(26018,G<br>lg= am                 | number= 4<br>replace(25686,A<br>/*tag= al              | number= 4<br>534029365<br>*tag= ak | standard name=<br>514825339<br>*tag= aj                | standard name=<br>eplace(23963,T<br>*tag= ai           | 'standard_name=<br>:eplace(22941,G<br>'*tag= ah        | standard_name=<br>eplace(22230,T<br>*tag=_aq           | standard_name=<br>eplace(21477,A<br>*tag= af           | standard_name=<br>eplace(21166,A<br>*tag= ae                     | <pre>etandard_name= eplace(20362,A *tag= ad</pre> | eplace(20204,G<br>*tag= ac             | ag= ab    |
| *tag= az                 | 37517<br>ay                 | eplace(30732,C)  *tag= ax  standard name= "Single nucleotide | standard name= "Single nucleotide<br>064030792<br>*tag= aw | /standard_name= "Single nucleotide<br>replace(29761,T)<br>/*tag= av | number= 5<br>number= 5<br>eplace(29572,T)<br>*tag= au | er = 86<br>er = 30639                                         | ar<br>dard_name= "Single nucleotide polymorphism"<br>29542 | <pre>/*tag= aq /standard name= "Single nucleotide replace(28772, A)</pre> | *tag= ap<br>standard_name= "Single nucleotide<br>eplace(28032,A) | U 10                                |                            | standard name= "Single nucleotide<br>eplace(26078,A)<br>*taq= an | - E                                                    | 4<br>25686,A)<br>al                                    | mber= 4<br>4029365<br>ag= ak       |                                                        | W 19                                                   | 0 11                                                   | W W                                                    | 6 5                                                    | standard_name= "Single nucleotide<br>eplace(21166,A)<br>*tag= ae | 6 5                                               | "Single                                | /*tag= ab |

```
RESULT 164
ADZ13149/c
ID ADZ131
XX ADZ1311
XX ADZ1311
XX DIAGNO
XX PLOMMAN
XX DIAGNO
XX DIAGNO
XX DIAGNO
XX DIAGNO
XX DOZOOS
XX WOO2005
XX WOO2005
XX JONAPR
XX 23-SEF
XX 23-SEF
XX CHIR
XX CHIR
XX WPI; 2
XX WPI; 2
XX WNuclei
PT compri
  맑
  S
  Query Match
Best Local Similarity
  Matches
  variation
  variation
  variation
   variation
  variation
   variation
   variation
          Nucleic acid array useful for detecting cancer associated nucleic acid, comprises two or more nucleic acid probes.
  WPI; 2005-273395/28
  Morris
  23-SEP-2003; 2003US-00669920
  23-SEP-2004; 2004WO-US031617
  07-APR-2005.
  WO2005031001-A2
   Homo sapiens
  cytostatic; gene;
   Diagnosis; DNA microarray; microarray; biochip; cancer; neoplasm;
  Human cancer-associated genomic DNA #56
   16-JUN-2005
  ADZ13149 standard; DNA; 57105
   variation
  variation
  variation
   (CHIR ) CHIRON CORP
  2898 GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 2942
  Ä
  GGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAG 25757
  Malandro
  Conservative
  (first entry
   replace(36681,G)
/*tag= hi
  /standard_name= "Single nucleotide
replace(34179,T)
  replace (36607,G)
  replace (34532,C)
   replace (34451, C)
   replace (34249,
   replace (32525,G)
  replace(32032,A)
  replace (31376, A)
  replace (36541,C)
   /standard_name= "Single nucleotide polymorphism
replace(34249,T)
   /standard_name= "Single nucleotide polymorphism"
  /standard_name= "Single
   standard_name= "Single
  standard_name=
   standard_name= "Single nucleotide"
  standard_name= "Single nucleotide polymorphism"
   standard_name= "Single nucleotide polymorphism"
  standard_name= "Single nucleotide polymorphism"
  standard name=
  *tag=
   100.0%;
  SK
  1.4%;
   ğ
   þ.
  Score 45;
Pred. No.
   0; Mismatches
   먪
   "Single
  "Single
  DB 12;
   2.8e-10;
  nucleotide polymorphism
   nucleotide polymorphism*
  nucleotide
   nucleotide
  <u>,,</u>
   Length 40090;
  Indels
  polymorphism"
   polymorphism"
   polymorphism
   polymorphism'
   Gaps
   0
  밁
   ક
   Best Loc
Matches
  Query Match
```

```
CC The invention also relates to a peptide array comprising two or more consolated polypeptides encoded by a CA nucleic acid sequence, a compound control that binds to a polypeptide, which is prepared by immunizing a host animal control that composition comprising the polypeptide or its fragment which composition comprising the polypeptide or its antigen binding control the antigen or its antigen binding fragment, a composition comprising the antibody and a carrier, a method of fereening for anticancer activity, a composition comprising the control of treating cancer and a method of inhibiting expression of a CA nucleic acid, a method of disgnosing cancer, a complete acid in a cell. The CA nucleic acids are useful for detecting CA concleic acids. The antibody is useful for detecting the presence or cancer cells in an individual which involves contacting cells from the individual with the antibody, where the detection of the complex correlates with the presence of cancer cells in the invention is also useful for dayneth of cancer cells in an individual. The invention is also useful for daynetic agent to cancer cells in an individual. The invention is also useful for daynession of a CA gene in the invention.

This sequence represents human cancer-associated genomic DNA of the coll. This sequence represents human cancer-associated genomic DNA of
Sequence 57105 BP; 15389 A; 12942 C; 12984 G; 15770 T; 0 U;
   The invention relates to a nucleic acid array for detecting associated (CA) nucleic acid, comprising two or more nucleic
  Disclosure; SEQ ID NO 669; 198pp; English
  invention.
   comprising two or more nucleic
   20
     20 Other;
   acid
   cancer
   probes
```

```
39001
                                 3078
                      GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   38957
                                  3122
```

Local Similarity nes 45; Conserv

Conservative 0;

1.4%;

Score 45; Pred. No.

DB 14; 1 . 2.8e-10;

Length 57105; Indels

0

Gaps

0

Mismatches

```
RESULT 165
ABK83563
ID ABK835
XX ABK835
XX ABK835
XX Human;
XX Human;
XX Human;
XX Human;
XX Human;
XX Human;
XX Freuman
XX Freuman
XX Grohn;
XX Grohn;
XX Grohn;
XX Homo s
XX Grohn;
XX Homo s
XX Homo s
XX Homo s
XX Grohn;
XX Homo s
XX H
  Human; ss; granulocytic cell; DNA chip; bacterial infection; viral infection; parasitic infection; protozoal infection; fungal infection; sterile inflammatory disease; psoriasis; rheumatoid arthritis; glomerulonephritis; asthma; thrombosis; cardiac reperfusion injury; renal reperfusion injury; ARDS;
  adult respiratory distress syndrome; inflammatory bowel disease; Crohn's disease; ulcerative colitis; periodontal disease; granulocyte activation; chronic inflammation; allergy.
  Human cDNA differentially expressed in granulocytic cells #134.
  ABK83563
  Homo sapiens.
   14-AUG-2002
   standard; cDNA; 57248
   (first entry)
```

WPI; 2002-435328/46

Beazer-Barclay Y, Weissman

Z.

Yamaga

ŝ

03-OCT-2000; 2000US-0237189P 03-OCT-2001; 2001WO-US030821.

GENE LOGIC INC.

11-APR-2002. WO200228999-A2.

```
PRACES
  밁
   Ş
  CC tissue, an altergic response in a subject exposure of a subject to a pathogen or sterile inflammatory disease, by detecting the level of cexpression in a sample of the tissue of gene(s) from Gs, where the level of expression of the gene is indicative of inflammation; (4) treating (MS) an inflammation (especially chronic) or in a tissue, an allergic response in a subject, exposure of a subject to a pathogen or sterile conflammatory disease, by contacting a tissue having inflammation with an cagent that modulates the expression of gene(s) from Gs in the tissue. M1 is useful for modulating GA, M3 is useful for screening an agent capable of modulating GA preferably in an C inflammation in a tissue; M4 is useful for detecting an inflammation (especially chronic) in a tissue, an allergic response in a subject, conflammation injury, renal reperfusion injury, axbs, adult creativatory distress syndrome, inflammatory bowel disease (e.g. C cardiac reperfusion injury, renal reperfusion injury, AxDs, adult creative, ulcerative colitis, periodontal disease, crohn's conditions. The present sequence represents a gene differentially conditions of the above conditions of the pinted specification, but was obtained in celectronic format directly from WIPO at C electronic format directly from WIPO at
   Matches
   Query Match
Best Local (
  The invention relates to detecting (M1) granulocyte (GC) activation (GCA), by detecting the level of expression of gene(s) (Gs) identified DNA chip analysis as given in the specification, and comparing the expression level to an expression level in an unactivated GC, where differential expression of Gs is indicative of GCA. Also included are modulating (M2) GA by contacting GC with an agent that alters the expression of at least one gene in Gs; (2) screening (M3) for an agent capable of modulating GCA or an inflammation (especially chronic) in a pathogen or sterile inflammatory disease using the gene expression profile; (3) detecting (M4) an inflammation (especially chronic) in a pathogen or allergic response in a subject, exposure of a subject to a pathogen and allergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease using the gene expression profile; (3) detecting (M4) an inflammation (especially chronic) in a response of a subject to 17-FEB-2004; 2004WO-US004730.
   02-SEP-2004
  WO2004074320-A2
  Human; ds; cancer-associated protein; gene; cytostatic; cancer;
leukaemia; lymphoma; CAP.
  Human cancer-associated genomic DNA HD18-038
  18-NOV-2004
  ABD32902
  ABD32902 standard; DNA; 65277
  Sequence 57248 BP; 15003 A; 13601 C; 13307
   Claim 1; SEQ ID NO 134; 114pp; English.
   Detecting granulocyte activation by detecting differential expression of genes associated with granulocyte activation, which serves as diagnostic markers that is useful for monitoring disease states and drug toxicity.
   2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 2932
   45
  Similarity
   TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 31345
   Conservative
  (first
  entry)
   1.4%;
  ٥,
  Score 45; ; Pred. No
  ₽₽
  Mismatches
  No.
  DB 6;
  2.8e-10;
  G; 15337 T; 0 U; 0 Other;
  Length 57248;
   (Gs) identified
  0
  Gaps
   in a
  λÂ
  0
```

```
CC mentioned nucleotide sequences, an isolated polypeptide (encoded within CC an open reading frame of a CA sequence selected from any of the 95 CC polynucleotide sequences as mentioned in the specification, or its CC complement), an isolated antibody, (or its antigen binding fragment) that binds to the above polypeptide, a hybridoma that produces the above CC monoclonal antibody, a pharmaceutical composition comprising the above CC antibody and a pharmaceutical excipient, a kit for detecting cancer CC cells (comprising the antibody cited above, methods for diagnosing cancer CC or for detecting the presence or absence of cancer cells in an CC individual, a method for inhibiting growth of cancer cells in an CC individual, a method for delivering a therapeutic agent to cancer cells in an individual, an electronic library comprising the above CC polynucleotide or polypeptide (or their fragments), methods of screening CC the activity of a CA protein (CAP), methods for detecting cancer CC associated with expression of a polypeptide in a test cell sample, a method for treating cancers and a method for inhibiting the expression of CC Agene in a cell. The composition and methods are useful for detecting, diameted and a method are useful for detecting.
  14-FEB-2003;
14-MAR-2003;
15-APR-2003;
15-JUN-2003;
13-JUN-2003;
   diagnosing, preventing and treating cancers, especially lymphoma and leukaemia. These may also be used in screening for agents that modul; cancer. The present sequence is a human CAP genomic sequence. Note: sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from W.
   The invention relates to an isolated nucleic acid comprising at least 10 contiguous nucleotides of any of the 233 polynucleotide sequences given in the specification, or its complement. The nucleic acids encode cancer associated proteins. Also included are an expression vector comprising the isolated nucleic acid cited above, a host cell comprising the above recombinant nucleic acid or expression vector, a microarray for detecting a cancer-associated (CA) nucleic acid comprising at least one probe comprising at least 10 contiguous nucleotides of any of the above-
  claim 16; seqid 602; 310pp; English.
   New isolated cancer-associated polynucleotides and polypeptides useful for diagnosing, preventing or treating cancers, especially lymphoma an leukemia, or in screening for agents that modulate cancer.
   Morris
   (SAGR-)
                                     ftp.wipo.int/pub/published_pct_sequences
   DW,
   SAGRES DISCOVERY INC.
   Morris DW,
   2003US-00417375.
2003US-00461862.
2003US-00663431.
  2003US-00737318
   2003US-00367094.
2003US-00388838.
   Malandro MS
  lymphoma and
  from WIPO
  detecting
   cancer-
```

```
RESULT 167
ACN43986/c
ID ACN439
   Matches
  Query Match
 18-NOV-2004
                           ACN43986;
   ACN43986 standard;
  Local Similarity
   45;
   GATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAGC 27364
   GATCACCTGAGGCCAGGAGTTCGAGACCAGCCTGGCCAACATAGC 2943
  Conservative
(first entry)
   DNA; 73995 BP
  100.0%;
   1.4%;
  0;
  Score 45;
Pred. No.
   Mismatches
   DB 13;
  2.8e-10;
   Length 65277;
   Indels
  0;
  Gaps
```

Human genomic sequence hCG40211.

밁 S

Sequence 65277 BP;

19651 A; 11706 C; 12664 G; 21256 T; 0 U; 0

Other;

```
RESULT 168
ACA64942
   S
  맑
   Query Match
Best Local :
  are associated with carcinomas. The sequences are useful for: (1) for screening drug candidates; (11) for screening of bloactive agent capable of binding to Carcinoma Associated Protein (CAP); (11) for screening of bloactive agent capable of modulating the activity of CAP; (1v) for a bloactive agent capable of modulating the activity of CAP; (1v) for evaluating the effect of a candidate carcinoma drug; (v) for diagnosing carcinoma; (vi) for inhibiting the activity of CAP; (11) for treating carcinoma; (vii) for nutralizing the effect of CAP; (1x) as a blochip; (x) for diagnosing carcinoma or a propensity to carcinoma; and (xi) for determining Carcinoma Associated (CA) gene copy number. In addition, the
  Human; chronic inflammatory joint disease; infection; tumour; antiinflammatory; cytostatic; antiarthritic; antirheumatic; immunosuppressive; gene therapy; etiological pathogenicity; ds
  determining Carcinoma Associated (CA) gene copy number. In addition, the CA genes are useful as DNA vaccines and the CAP are useful as markers of carcinoma including lymphoma. The present sequence is one such CA coding sequence. Note: This patent is an equivalent to basic patent responsive seasons.
  The present invention are associated with
   01-MAR-2002; 2002US-00087192.
  Cytostatic; carcinoma; lymphoma; cancer; human; gene; ss
  ACA64942;
   Sequence 73995 BP; 17594 A; 18402 C; 19247 G; 18752 T; 0 U; 0 Other;
   sequence. Note:
US2002182586A1,
  Claim 1; SEQ ID NO 208; Opp; English.
   comprises a
  Recombinant nucleic acid useful for diagnosis and treatment of carcinoma
   28-FEB-2003; 2003WO-US006235
   12-SEP-2003.
   WO2003073826-A2
  Homo sapiens
30-MAY-2001; 2001DE-01027572
  Homo
  Human FRAP1 DNA corresponding to AL049659.
  27-JUN-2003
   ACA64942 standard; DNA; 78539
                                  30-MAY-2001; 2001DE-01027572.
  immunosuppressive;
   (SAGR-) SAGRES DISCOVERY
  27485 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 27441
   2003-328604/31.
  sapiens.
   45;
   Similarity
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  invention relates to
   nucleotide sequence.
   Conservative
  (first
  for which no sequence data was published
  entry)
   1.4%; Score 45;
100.0%; Pred. No.
   ..
   Mismatches
  novel DNA and protein sequences which
   DB 11;
2.8e-10;
   0
   Length 73995;
   Indels
   0,
   Gaps
   0
```

```
RESULT 169
ADD79404/c
ID ADD79404;
XX
AC ADD79404;
XX
DT 26-AUG-2004 (first entry)
XX
DE DPF3 region, SEQ ID 3.
XX
CYCOStatic; Gene therapy; breast cancelled the control of the control o
  S
   밁
  Best Local Similarity
Matches 45; Conserv
   Query Match
  products of the invention have antiinflammatory, cytostatic, products of the invention have antiinflammatory, cytostatic, antirheumatic and immunosuppressive activity and can be used for gene therapy. The reagent of the invention and any proteins and antibodies derived from it, are used (i) for analysing tissue and blood samples for medical diagnosis; (ii) for diagnosis and characterisation of chronic joint disease, on the basis of molecular characterisation, and determining the etiological pathogenicity principle of as yet uncharacterised inflammatory diseases, also monitoring progression and/or treatment of disease, and optimisation of therapy and (iii) for developing treatments for inflammatory diseases, particularly of joints, infections and tumours. ACA64801-ACA64965 represent human polynucleotides used in the method of the invention
  This invention describes a novel reagent for diagnosis, molecular definition and therapy of chronic inflammatory joint diseases, and other inflammatory disorders, infective or tumour diseases in humans. The
  Cytostatic; Gene therapy; breast cancer; human; DLG1; KIAA0783; DPP3; CENPC1; gene; ds; SNP; single nucleotide polymorphism; D4, zinc and double PHD fingers, family 3; CERD4; cer-d4; PLJ14079; D4, 2100 SNR; Rho family guanine-nucleotide exchange factor;
   Sequence 78539 BP; 23554 A; 17605 C; 17140 G; 20240 T; 0 U; 0
  Claim 1; Page; 12pp; German
   proteins.
   Reagents for diagnosis, study and therapy of chronic inflammatory joint and other diseases, comprises any of many specified genes or derived
   WPI; 2003-240797/24.
   (PATH-) PATHOARRAY GMBH
  56640
  3078
   H
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  GTGCCACTGCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  Conservative
   Location/Qualifiers
   /standard_name= "Single nucleotide polymorphism"
/note= "This SNP is described as a A/G SNP"
  /*tag= a
/standard name= "Single nucleotide polymorphism"
/note= "This SNP is described as a A/C SNP"
   /standard name= "Single nucleotide polymorphism"
/note= "This SNP is described as a T/G SNP"
   1.4%;
   ď
   Blaess
   0,
  Score 45;
Pred. No.
   Mismatches
   S
  DB 8; Lo
   0
   Length 78539;
   Indels
  56684
   0
   Other;
  Gape
   0
```

| /note= "This SNP is described as a G/C SNP" 37314 /*tag= v /standard_name= "Single nucleotide polymorphis" | /standard name= "Single nucleotide<br>/note= "This SNP is described as a<br>variation 36254<br>/*tag= u | /*tag=<br>/standar<br>/note= "<br>variation 35856                                                                                  | variation 35588  /*tag= r /standard name= "Single nucleotide /note= "This SNP is described as a variation 35619 | /note= "This SNP is described as a T/C SNP" 32003 /*tag= q /standard name= "Single nucleotide polymorphi /note= "This SNP is described as a A/C SNP" | /*tag= 0 /standard name= "Single nucleotide /note= "This SNP is described as a variation /*tag= p /*tag= p /standard name= "Single nucleotide | variation variation                                                                                       | variati                        | variation 2468 /*ta // gta // not | variation /note= "This SNP /4582 k /standard name= /note= "This SNP /note= | /*tag= i /standard name= "Single nucleotide polymorphi /note= "This SNP is described as a A/G SNP" variation 18858 | variation 1865 /*ta /sta /not variation 1869                                                    | ria ti | variation                                                                                               | FT /standard name= "Single nucleotide polymorphism" FT /note= "This SNP is described as a T/C SNP" |
|------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------|--------------------------------|-----------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------|--------|---------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------|
| variation 89751 /*tag= an /standard name= "Single nucleotide /note= "This SNP is described as a            | /note= "This SNP is decoration 85507  /*tag= am /*standard name= "Single /note= "This SNP is dec        | /standard name= "Single nucleotide /note= "This SNP is described as a variation 72752 /*tag= al /standard name= "Single nucleotide | variation /                                                                                                     | variation 60682  /*tag= ai /standard name= "Single nucleotide /note= "This SNP is described as a variation 61291                                     | variation 5                                                                                                                                   | /*tag= at /standard_name= "Single nucleotide /note= "This SNP is described as a variation 53971 /*tag= ag | /*tag=<br>/standan<br>/note= ' | /*tag=<br>/standar<br>/note= "    | variation 49774  /*tag= ac /standard name= "Single nucleotide /note= "This SNP is described as a variation 51796                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               | variation 4                                                                                                        | /*tag= z<br>/standard name= "Single<br>/note= "This SNP is deal<br>variation 43090<br>/*tag= aa | Hoy H  | /*tag= w "Single nucleotide /standard name= "Single nucleotide /note= "This SNP is described as a 40095 | variation                                                                                          |

```
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XX ABD335
XX ABD335
XX Haman;
XW Human;
XW Human;
XW Homo &
YX WO2004
XX Homo &
YX Ho
   RESULT 170
   밁
  र्
  Query Match
Best Local Sim
Matches 45;
    The invention relates to cancer-associated proteins (CAP) and the cancer-associated (CA) nucleic acids encoding them. The invention also relates to a method for treating cancers involving administering to a patient an inhibitor of CAP, and a method of screening for anticancer activity in a potential drug involving providing a cell that expresses a CA gene, contacting a tissue sample derived from a cancer cell with an anticancer
  Identifying a subject at risk of of polymorphic variations in the which are associated with breast
   25-NOV-2002; 2002US-0429136P.
24-JUL-2003; 2003US-0490234P.
  Novel human cancer associated protein encoded within open reading f of cancer associated gene, useful as targets for diagnosing cancer.
   Human; cancer-associated protein; CAP; cancer-associated gene;
  Human cancer-associated (CA) gene HD07-103
   Claim 24; Fig
   Roth RB,
   25-NOV-2003; 2003WO-US037943
  10-JUN-2004.
  WO2004047514-A2.
  Morris
  15-JUL-2004.
  WO2004058146-A2
   Homo sapiens
   ds; cancer; cycostatic
   18-NOV-2004
  ABD33524 standard; DNA; 107543 BP
  (SAGR-)
   17-DEC-2002; 2002US-00322281
   15-DEC-2003; 2003WO-US040081.
  Local Similarity
   59096
   3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   2004-441037/41.
  16;
  Ä
  SAGRES DISCOVERY INC
  SEQUENOM
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 59052
   Nelson MR,
  SEQ ID NO
  Malandro
  Conservative
   (first entry)
   u
T
  INC
   227pp; English
  1.4%; STEEL NO. 100.0%; Pred. 
   706; 182pp;
  S
  Braun
   Þ
  Kammerer SM,
  English.
  breast cancer by detecting the presence DLG1, KIAA0783, DPF3 or CENPC1 regions cancer in a nucleic acid sample from a
   DB 12;
2.7e-10;
0;
  Reneland
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RESULT 171
ABD33242/c
ID ABD33242/c
AC ABD3322
XX ABD332
XX ABD332
XX Human
XX Human;
XW Human;
XW Homo s
PN WO2004
XX IS-JUL

   밁
  S
  The invention relates to cancer-associated proteins (CAP) and the cancer-
CC associated (CA) nucleic acids encoding them. The invention also relates
CC to a method for treating cancers involving administering to a patient an
CC inhibitor of CAP, and a method of screening for anticancer activity in a
CC contacting a tissue sample derived from a cancer cell with an anticancer
CC contacting a tissue sample derived from a cancer cell with an anticancer
CC contacting at monitoring the effect of the anticancer drug candidate and monitoring the effect of the anticancer drug candidate and monitoring the effect of the anticancer drug candidate
CC on expression of the CA gene. The CAP proteins are useful for detecting
CC cancer associated with expression of a CAP protein in a test cell sample
CC and for screening for a bioactive agent capable of modulating the
CC activity of a CAP protein. The CA nucleic acids are useful for diagnosing
CC cancer, involving determining the expression of a CA nucleic acid in a
CC tissue. This sequence represents a human CA gene of the invention. Note:
CC specification, but was obtained in electronic format directly from WIPO
CC at for the capter of the capter of the printed
CC specification, but was obtained in electronic format directly from WIPO
   Matches
  Query Match
Best Local Similarity
   drug candidate and monitoring the effect of the anticancer drug candidate on expression of the CA gene. The CAP proteins are useful for detecting cancer associated with expression of a CAP protein in a test cell sample and for screening for a bloactive agent capable of modulating the activity of a CAP protein. The CA nucleic acids are useful for diagnosing cancer, involving determining the expression of a CA nucleic acid in a tissue. This sequence represents a human CA gene of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
   Novel human cancer associated protein encoded within open reading frame of cancer associated gene, useful as targets for diagnosing cancer.
   WPI; 2004-499109/47
   Homo sapiens
  Human; cancer-associated protein; CAP; cancer-associated gene;
  18-NOV-2004
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   Morris
   17-DEC-2002; 2002US-00322281.
   15-DEC-2003; 2003WO-US040081.
   15-JUL-2004.
   (SAGR-)
   ftp.wipo.int/pub/published_pct_sequences
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  cancer; cytostatic
   3078
   16; SEQ ID NO 268; 182pp; English.
   cancer-associated
   Ð₩,
   SAGRES DISCOVERY INC
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   Malandro
   Conservative
  (first entry)
  1.4%;
   š
   ξ
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   0
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Pred. No.
  gene HD07-040
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  2.7e-10;
   0
   Length 107543;
   Indels
   3122
  Ş
   Gaps
   0,
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ftp.wipo.int/pub/published\_pct\_sequences

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RESULT 173

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WP ABXO8336_01
WP ABXO8336_03
WP ABXO8336_03
WP ABXO8336_06
WP ABXO8336_06
WP ABXO8336_07
WP ABXO8336_07
WP ABXO8336_07
WP ABXO8336_07
WP ABXO8336_09
WP ABXO8336_10
WP ABXO8336_11
AAD53224_2/c
AAD53224_2/c
Continuation (3 of 6)
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WP AAD53224_2
WP AAD53224_3
WP AAD53224_3
WP AAD53224_4
WP AAD53224_5
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ABL57909 2
COntinuation
WP Sequence spragmen
WP Fragmen
WP ABL579
WP ABL579
WP ABL579
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   5
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   ÖS
   Query Match
Best Local S
Matches 45
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P Fragment Name
P ABL57909 0
ABL57909 1
ABL57909 2
ABL57909 3
  Query Match
Best Local S
Matches 45
  Query Match
Best Local S
Matches 45
   Sequence
   21081
   61071
   52355
   3078
  3078
  l Similarity
45; Conser
  1 Similarity
45; Conserv
   Similarity
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  107745
   of 6) c
  1.4%;
larity 100.0%;
Conservative
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   17
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11300001
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   ; Score 45; DB
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  rom base
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End
110000
210000
310000
368004
   22736
  from
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110000
210000
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410000
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  Shoot
   FOCUS
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   base 200001
TUS AAD53224
  Ç
   base 300001
CUS ABX08336
   nse 200001
NBL57909
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  DB 6;
3. 2.7e-10;
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   (Human chromosome Accession Aad53224
  (Human transporter Accession Abl57909
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   (Human phosphodiesterase Accession Abx08336
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   Length 107745;
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   Indels
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  52399
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  w
  Gaps
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  protein
   Other;
   q-arm
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  breakpoint
  0
  0
  0
   gene
   (PDE4D)
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   片
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RESULT 176

ADN97989_03/c

ADN97989_03/c

Continuation (4 of 17)

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WP ADN97989_03

WP ADN97989_05

WP ADN97989_05

WP ADN97989_06

WP ADN97989_07

WP ADN97989_07

WP ADN97989_08

WP ADN97989_10

WP ADN97989_11

WP ADN97989_13

WP ADN97989_13

WP ADN97989_13

WP ADN97989_13

WP ADN97989_15

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WP ADJ2598
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ADJ2598
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Best Local S
Matches 45
  Query Match
Best Local S
Matches 45
   P Sequence split into P Fragment Name P Fragment Name P ADJ25985 00 P ADJ25985 03 P ADJ25985 06 P ADJ25985 06 P ADJ25985 06 P ADJ25985 06 P ADJ25985 07 P ADJ25985 10 P ADJ25985 11 P ADJ25985 12 P ADJ25985 13 P ADJ25985 14 P ADJ25985 15 ADJ25985 15 ADJ25985 15 ADJ25985 16
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 61071
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   3078
  3078
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   1 Similarity
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GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
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  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
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  1.4%;
ilarity 100.0%;
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llarity 100.0%; I
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11000001
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400000
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  ADN97989 from fragments LO
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  Score 45; DB; Pred. No. 2.7
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910000
9101000
1110000
1310000
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1510000
1691138
   Bnd
11000
210000
310000
410000
510000
610000
610000
910000
910000
910000
1110000
1210000
1310000
1410000
1510000
1610000
  rom base 300001
LOCUS ADN97989
  LOCUS
  Mismatches
   base 300001
CUS ADJ25985
   2.
2.
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   DB 8; 1
2.7e-10
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  Length
   Length 110000;
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   Indels
   Indels
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 61027
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  3122
   61026
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ARBSULT 178
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Continuation (continuation)
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WP Sequence sp
WP ARBS5189
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ADO50281_03/c
Continuation (
  SHARAB
   Ś
  맑
  S
  RESULT 179
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  Query Match
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   Matches
   Matches 45;
   Query Match
   ntinuation (4 of 17)
Sequence split into
   Sequence split into
  Fragment Name
Fragment Name
AEB85185 00
AEB85185 02
AEB85185 03
AEB85185 04
AEB85185 06
AEB85185 06
AEB85185 07
AEB85185 09
AEB85185 10
AEB85185 11
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AEB85185 11
AEB85185 13
AEB85185 13
AEB85185 14
AEB85185 15
         26-AUG-2004
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AD050281 02
AD050281 02
AD050281 03
AD050281 04
AD050281 04
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AD050281 06
AD050281 07
AD050281 07
AD050281 10
AD050281 11
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   61070
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   Similarity
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 61027
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
  ilarity 100.0%; 1 Conservative 0;
   Conservative
  into
        (first entry)
   17
17
   of
17
  DNA;
   1400001
1500001
1600001
  1400001
1500001
1600001
   800001
900001
1000001
1100001
1200001
   1100001
  1000001
   1300001
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   1300001
  AEB85185 from base 300001 (Human phosphodiesterase 4D fragments LOCUS AEB85185 Accession Aeb85185
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500001
  100001
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   500001
  700001
  100001
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  300001
  600001
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Pred. No.
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Pred. No.
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1110000
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710000
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1410000
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310000
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   2.7e-10;
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  Length 110000;
  Indels
   Indels
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   0
   0
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  SE(
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ABT10719

ABT10719

standard; cDNA;

122748

ΒP

01-AUG-2002.

WO200259271-A2

25-JAN-2002; 2002WO-US002176.

cytostatic;

gene

therapy; gene; ss.

Human; breast specific gene; breast cancer; differential expression;

Human breast cancer associated coding sequence SEQ ID NO:

04-DEC-2002

(first

entry)

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밁
  á
RESULT 180
   Query Match
Best Local &
  Matches
  which comprises obtaining a first soft tissue sample from an individual and a normal soft tissue sample from the same or different individual, determining the expression of a gene in both samples and comparing the expression of the gene in both soft tissue samples, where a higher level of protein expression in the first soft tissue sample indicates the presence of soft tissue sarcoma. The method of the invention has cycostatic applications and may be useful for detecting soft tissue sarcoma, possibly via gene therapy or vaccine production. The nucleic acid sequences may be useful in diagnostic and screening applications. The current sequence is that of a human soft tissue sarcoma-upregulated DNA of the invention. The current sequence is not shown within the
   Early detection of soft tissue sarcoma comprises determining expression of a gene in a first soft tissue sample and a normal soft tissue sample and comparing the gene expression, also useful in treating soft tissue
  Sequence 116561 BP; 32234 A; 27837 C; 28253 G; 28237 T; 0 U; 0 Other.
   The invention relates to a novel method for detecting soft tissue sarcoma which comprises obtaining a first soft tissue sample from an individual
   and comparing the gene
   Aziz N,
   26-NOV-2002; 2002US-0429739P
  10-JUN-2004.
  WO2004048938-A2
   Human soft tissue sarcoma-upregulated DNA - SEQ ID 409
  specification per se but was submitted in CD format by the inventor.
  Example 2;
   WPI; 2004-441208/41
  26-NOV-2003; 2003WO-US038193
  (PROT-)
  104267
   3078 GTGCCACTGCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   tissue sarcoma; cytostatic; gene therapy; vaccine; screening;
  l Similarity
45; Conserv
  PROTEIN DESIGN LABS INC.
   Ginsburg WM,
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 104311
  SEQ ID NO 409; 210pp; English
  Conservative
   1.4%; 5-
100.0%; Prr
   Zlotník A;
   Score 45;
Pred. No.
  Mismatches
   DB 12; I
2.7e-10;
  Length 116561;
  Indels
  3122
  0
  Gaps
   human;
  0
```

```
RESULT 181
ADL13909/c
ID ADL139
XX ADL139
XX ADL139
XX OSTEODA
XX HOMO 8
XX HOMO 8
XX HOMO 8
XX I9-DEC
XX 19-DEC
XX INCY-
XX JONES
XX JONES
XX JONES
XX JONES
XX JONES
XX JONES
   밁
  S
  Query Match
Best Local
   Matches
   patient, which comprise detecting the level of expression in a tissue sample of two or more genes selected from those shown in ABT09867-ABT1112, where a differential expression of the genes indicates breast cancer. The methods are useful in diagnosing, treating, detecting the progression, and in monitoring treatment of breast cancer in patients. The methods are also useful as a screening tool for agents that modulate the onset or progression of breast cancer. The breast cancer genes may be used as diagnostic markers for the prediction or identification of the malignant state of breast tissue, for confirming the type and progression of cancer, and for drug screening and assays. The present sequence is a coding sequence of the invention. Note: The sequence data for this patent
  ds; gene; osteopathic; antiinflammatory; antiarthritic; ge joint space narrowing; osteophyte development; joint pain; osteoarthritis; SNP; single nucleotide polymorphism.
   Sequence 122748 BP; 32088 A;
  did not form part of the printed specification, but was obtained electronic format directly from WIPO at
  Diagnosing breast cancer in a patient comprises detecting the level gene expression in cell or tissue samples, where a differential gene expression is indicative of breast cancer.
  Orr MS,
  25-JAN-2001; 2001US-0263757P.
25-APR-2001; 2001US-0286090P.
23-MAY-2001; 2001US-0292517P.
 WPI; 2003-559141/52
                                  Jones KA,
   19-DEC-2002; 2002WO-US041225
  WO2003054166-A2
   Osteoarthritis-associated polymorphic nucleotide #441.
  06-MAY-2004 (first entry)
   ADL13909
   ADL13909 standard; DNA; 129588 BP
  ftp.wipo.int/pub.published_pct_sequences
   Claim 1;
   The present invention relates to methods of diagnosing breast cancer
  20-DEC-2001; 2001US-0342603P
   (GENE-)
   (INCY-) INCYTE GENOMICS
   37876
   sapiens
   2002-674803/72.
   Similarity
   GENE LOGIC INC
   GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 37920
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
   SEQ ID NO 853; 260pp + Sequence Listing; English
  Nation M,
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  100.0%;
  1.48;
   0;
  Score 45;
Pred. No.
   31056 C; 30547 G;
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  DB 6; Le 2.7e-10;
   0,
  Length 122748;
   29057 T; 0 U; 0 Other;
   Indels
   gene therapy;
   0;
   Gaps
  gene
  'n
   얁
  in a
   0,
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밁 5

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XFFFFX8X555555555555555555555
  Matches 45;
                         Query Match
Best Local
   The invention relates to a method of determining susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain comprising identifying whether the individual has at least one polymorphism in a polymuclectide encoding at least one of the protein listed in the specification. The methods, composition and agent are useful for modulating the susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain that is associated with a disease, preferably osteoarthritis. The cell line and the non-human animal are useful for screening for an agent for diagnosing an individual having susceptibility to joint space narrowing and/or osteophyte development and/or joint pain. This sequence corresponds to the polymucleotide encoding a protein listed in the specification. (Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from WIPO at
   Determining susceptibility of an individual to joint space narrowing, osteophyte development and/or joint pain comprises identifying whether the individual has at least one polymorphism in a polynucleotide encoding
   Sequence 129588 BP; 35710 A; 27530 C; 26424 G;
   Disclosure; SEQ ID NO 441;
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   ftp.wipo.int/pub/published_pct_sequences).
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                     1.4%;
   297pp; English
0;
                       Score 45;
Pred. No.
  Mismatches
  DB 10;
                       2.7e-10;
  0;
   36620 T; 0 U; 3304 Other;
  Length 129588;
  Indels
  0;
  Gaps
  0
```

```
RESULT 182
ABK84797/c
   viral infection; parasitic infection; protozoal infection; fungal infection; sterile inflammatory disease; psoriasis; rheumatoid arthritis; glomerulonephritis; asthma; thrombosis;
   Crohn's disease; ulcerative colitis; periodon granulocyte activation; chronic inflammation;
  cardiac reperfusion
   Human; 88; granulocytic cell; DNA chip; bacterial infection;
   ABK84797
WO200228999-A2
  Homo sapiens.
   adult respiratory distress syndrome;
   Human cDNA
  27247
  3073 AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAGAGCAGAGCTCT 3117
  standard; cDNA; 149671
  AGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCT 27203
   differentially expressed in granulocytic cells #1368.
   (first entry)
  injury; renal reperfusion injury; ARDS; istress syndrome; inflammatory bowel disease;
  periodontal disease;
  allergy
```

Detecting granulocyte activation by detecting differential expression of genes associated with granulocyte activation, which serves as diagnostic markers that is useful for monitoring disease states and drug toxicity.

WPI; 2002-435328/46.

Beazer-Barclay Y, Weissman

SM,

Yamaga

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Vockley

(GENB-) GENE LOGIC INC.

03-OCT-2001; 2001WO-US030821.

2000US-0237189P

11-APR-2002

Claim

1; SEQ ID

NO 1368; 114pp; English.

```
RESULT 183
ADB70361/c
ID ADB703
XX ADB703
XX ADB703
XX ADB703
XX Moesin
XX Moesin
XX Gancer
KW Guamo
KW Homo s
XW Homo s
XX Homo s
XX PN WO2003
XX PP 05-SEF
XX
  닭
  Ś
  expression in a sample of the tissue of gene(s) from Gs, where the level CC of expression of the gene is indicative of inflammation; (4) treating CC (M5) an inflammation (especially chronic) or in a tissue, an allergic cresponse in a subject, exposure of a subject to a pathogen or sterile conflammatory disease, by contacting a tissue having inflammation with an eagent that modulates the expression of gene(s) from Gs in the tissue. M1 is useful for screening an agent capable of modulating GA, preferably in an inflammation in a tissue, M4 is useful for modulating GA, M3 is useful for screening an agent capable of modulating GA preferably in an inflammation in a tissue, M4 is useful for detecting an inflammation CC inflammation in a tissue, M4 is useful for detecting an inflammation CC exposure of a subject to a pathogen or sterile inflammatory disease (e.g. CC paoriasis, rheumatoid arthritis, glomerulonephritis, asthma, thrombosis, CC cardiac reperfusion injury, renal reperfusion injury, ARDS, adult CC expressed in geramicory parasitic infection, protozoal infection, infection, parasitic infection, protozoal infection, viral infection, parasitic infection, protozoal infection, parasitic infection, protozoal infection, parasitic infection, parasitic infection, parasitic infection, parasitic infection, parasitic infection, protozoal infection, the present sequence represents a gene differentially conform part of the printed specification, but was obtained in the conformation of the printed specification, but was obtained in
  Query Match
Best Local :
   modulating (M2) GA by contacting GC with an agent that alters the expression of at least one gene in Gs; (2) screening (M3) for an agent capable of modulating GCA or an inflammation (especially chronic) in a tissue, an altergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease using the gene expression profile; (3) detecting (M4) an inflammation (especially chronic) in a tissue, an altergic response in a subject, exposure of a subject to a pathogen or sterile inflammatory disease, by detecting the level of
   cancer; malignant pleural mesothelioma; MPM; lung adenocarcinoma;
squamous carcinoma; medulloblastoma; prostate cancer; breast cancer;
diffuse large B-cell lymphoma; follicular lymphoma; ovarian cancer;
   DNA chip analysis as given in the specification, and comparing the expression level to an expression level in an unactivated GC, where differential expression of Gs is indicative of GCA. Also included are modulating (M2) GA by contacting GC with an agent that alters the
   04-DBC-2003
   Sequence 149671 BP; 45600 A; 33308 C; 32389 G; 38374 T; 0 U; 0
                         05-SEP-2002; 2002WO-US028203
   13-MAR-2003.
  WO2003021229-A2
  Homo sapiens
  Moesin
   ADB70361;
  ADB70361 standard; cDNA; 149671 BP
   ftp.wipo.int/pub/published_pct_sequences
   Local Similarity
  94008 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 93964
   invention relates to
   2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 2932
   45;
  CDNA
  by detecting the level of expression of gene(s)
   SEQ ID
   Conservative
   (first entry)
   NO:53
   100.0%;
   1.4%;
   detecting (M1) granulocyte (GC)
   Score 45; ; Pred. No.
   <u>,</u>
   Mismatches
   DB 6; I
2.7e-10;
   ..
  Length 149671;
   Indels
   ଜୁ
   activation
  identified by
   0
  Other
   0
```

```
RESULT 184
ADJ371140/c
ID ADJ371
XX ADJ371
XX ADJ371
XX Pluman
XX Human;
XW Human;
XW Lung a
XW Follic
XX Folloo 8
XX US2003
XX US2003
XX PD 27-NOV
XX PF 05-SEF
PR 05-SEF
PR 30-AUC
XX PA (BGHM
XX (BGHM
XX (BDJ371)
  밁
   Ş
  Matches
   Best
  Query Match
   of genes that are differentially expressed in cancerous or non-cancerous conditions, determining the expression levels of the set of genes and calculating a ratio of the expression levels of the differentially expressed genes. M is useful for diagnosing the presence of cancer cells or non-cancer cells in a tissue sample, where the cancer is malignant pleural mesothelioma (MPM), lung adenocarcinoma, squamous carcinoma, medulloblastoma, prostate cancer, breast cancer, diffuse large B-cell lymphoma, follicular lymphoma and ovarian cancer, and for determining prognosis or outcome of a cancer patient. The ratio of expression levels of differentially expressed genes is used as an indicator of cancer type, cancer canse, and/or cancer prognosis, all of which are useful for determining a course of treatment of a patient. The present sequence encodes a human protein which is used in an example from the present
   The present invention describes a method (MI) for diagnosing the presence of cancer cells or non-cancer cells in a tissue sample, or determining the prognosis or outcome of a cancer patient. MI involves providing a set
   Diagnosing cancer cells in tissue sample, or determining prognosis or outcome of cancer patient, by calculating ratio of expression levels of genes that are differentially expressed in cancer and non cancer tissues.
   05-SEP-2001; 2001US-0317389P.
30-AUG-2002; 2002US-0407431P.
  Homo sapiens
  Human; malignant pleural mesothelioma; MPM; gene; ss; tumour; lung adenocarcinoma; squamous carcinoma; medulloblastoma; prostate cancer; breast cancer; diffuse large B-cell lymphoma;
   Human malignant pleural mesothelioma (MPM) cDNA #23
   ADJ37140 standard; cDNA; 149671 BP
   Sequence 149671 BP;
  Claim 67; Page 181-263; 396pp; English.
   Gordon GJ,
  05-SEP-2001; 2001US-0317389P.
30-AUG-2002; 2002US-00236031.
  05-SEP-2002; 2002US-00236031.
  27-NOV-2003
   US2003219760-A1
  22-APR-2004
  follicular lymphoma;
   Local
  94008
   2888
  2003-290233/28
  1 Similarity
45; Conserv
   BRIGHAM & WOMENS HOSPITAL INC
   TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG
  TGAGGCAGGTGGATCACCTGAGGCCCAGGAGTTCGAGACCAGCCTG 93964
   Jensen RV, Gullans SR, Bueno
  invention describes a method (MI) for diagnosing the presence
  Conservative
  (first entry)
  100.0%; F1
   45600 A;
  ovarian
   Score 45; pred. No.
   33308 C; 32389 G;
  Mismatches
   DB 9; Le
2.7e-10;
   in cancer and non cancer tissues.
  0,
  Length 149671;
   38374 T; 0 U; 0 Other;
  Indels
   2932
  0
  levels of
  Gaps
```

0

(BGHM ) BRIGHAM & WOMENS HOSPITAL INC

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   cc cells or non-cancer cells in a tissue sample, determining prognosis or contcome of a cancer patient, selecting a course of treatment for a cc subject having or suspected of having malignant pleural mesotheliona (MPM) and evaluating treatment of MPM comprising determining the ratio of c the expression level of a set of genes differentially expressed in a cc cancer tissue. The cancer is chosen from MPM, lung adenocarcinoma, c diffuse large B-cell lymphoma, follicular lymphoma and ovarian cancer. CC diffuse large B-cell lymphoma, follicular lymphoma and ovarian cancer. CC The method is useful for diagnosing MPM in a subject suspected of having CC MPM which involves obtaining a tissue sample suspected of being cancerous from a subject and determining the expression of nucleic acid markers or c its expression products in the tissue sample. This sequence are presents cuman MPM cDNA of the invention. Note: The sequence data for this patient CC did not form part of the printed specification but was obtained in clectronic format directly from USPTO at sequence absolute.html.
  Query Match
Best Local S
Matches 45
Disclosure; SEQ ID NO 879; 198pp; English
                                 Nucleic acid array useful for detecting cancer comprises two or more nucleic acid probes.
   07-APR-2005
   Human cancer-associated genomic DNA #75.
   16-JUN-2005
  Sequence 149671 BP; 45600 A; 33308 C; 32389 G; 38374 T; 0 U; 0 Other;
  Claim 44; SEQ ID NO 53; 53pp; English.
  determining ratio of expression cancer tissues.
  Diagnosing the presence of cancer or non-cancer cells in tissue sample, useful for diagnosing malignant pleural mesothelioma comprises determining ratio of expression level of a set of genes expressed in
   P-PSDB; ADJ37141
   Gordon
   23-SEP-2004; 2004WO-US031617.
   WO2005031001-A2
   Homo sapiens
   cytostatic; gene;
   Diagnosis; DNA microarray; microarray; biochip; cancer; neoplasm,
  ADZ13359 standard; DNA; 171398 BP
   Morris
   23-SEP-2003; 2003US-00669920
   (CHIR )
  Local Similarity
  94008
  invention relates to
   2888 TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 2932
   2005-273395/28
   2004-141744/14.
   <u>ج</u>
   DW, Malandro MS
   CHIRON CORP
  TGAGGCAGGTGGATCACCTGAGGCCAGGAGTTCGAGACCAGCCTG 93964
  Conservative
   Jensen
   (first entry)
   80
   R۷,
  100.0%;
   1.4%; Score 45;
100.0%; Pred. No.
   Gullans
  a method of diagnosing the presence of cancer
   0;
  Mismatches
   SR,
   Bueno
   DB 12; I
. 2.7e-10;
   associated
   Length 149671;
  Indels
   nucleic
  0
  Gaps
   acid
  0
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≵2222222222222222222222222223
  CC with a composition comprising the polypeptide or its antigen binding cells from the host expressing antibodies against the antigen or its antigen binding fragment, a composition comprising the cantibody and a carrier, a method of screening for anticancer activity, a composition comprising the cell of detecting a CA nucleic acid, a method of diagnosing cancer, a comprising cancer, a composition comprising the nucleic acid in a cell. The CA nucleic acids are useful for detecting CA concleic acids. The antibody is useful for detecting the presence or complex correlates with the antibody, where the detection cells from the individual with the antibody, and detecting the presence of complex correlates with the presence of cancer cells in an individual. The invention is useful for indignosing cancer cells in an individual. The invention is also useful for diagnosing cancer, for treating cancer and for inhibiting expression of a CA gene in a cell. This sequence represents human cancer-associated genomic DNA of
   The invention relates to a nucleic acid array for detecting a cancer associated (CA) nucleic acid, comprising two or more nucleic acid probes. The invention also relates to a peptide array comprising two or more isolated polypeptides encoded by a CA nucleic acid sequence, a compound that binds to a polypeptide, an isolated antibody or its fragment which binds to a polypeptide, which is prepared by immunizing a host animal
Sequence 171398 BP; 51304 A; 33907 C; 34019 G;
  the invention.
  52168 T;
0 U; 0 Other;
  probes.
```

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Matches
   Query Match
  Local
84806
                            3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACT 3115
  1 Similarity
45; Conserv
CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACT 84850
  1.4%; Score 45; DB llarity 100.0%; Pred. No. 2. Conservative 0; Mismatches
   DB 14;
   2.7e-10;
hes 0;
  Length 171398;
   Indels
   0
   Gaps
   0
```

```
RESULT 186
   ds; gene; osteopathic; antiinflammatory; antiarthritic;
   Osteoarthritis-associated polymorphic nucleotide #312.
  06-MAY-2004 (first entry
   ADL13780 standard; DNA; 190117 BP
```

joint space narrowing; osteophyte development; joint osteoarthritis; SNP; single nucleotide polymorphism. Homo sapiens rtnritic; gene therapy; joint pain;

WO2003054166-A2

19-DEC-2002; 2002WO-US041225

20-DEC-2001; 2001US-0342603P

(INCY-) INCYTE GENOMICS

Jones KA, Schafer A;

WPI; 2003-559141/52.

Determining susceptibility of an individual to joint space narrowing, osteophyte development and/or joint pain comprises identifying whether the individual has at least one polymorphism in a polymucleotide encoding

Disclosure; SEQ ID NO 312; 297pp; English

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```
RESULT 187
ABD33586
ID ABD335
XX ABD335
XX ABD335
XX Human
XX Human;
XW da; ca
XX Homo a
CX Homo a
CX Homo a
CX CAGR-
PT Of Calim
XX Homo a
CX The il
CC Lonia
CC Lonia
CC Contax
CC Conta
   Query Match
Best Local
   Matches
The invention relates to cancer-associated proteins (CAP) and the cancer-associated (CA) nucleic acids encoding them. The invention also relates to a method for treating cancers involving administering to a patient an inhibitor of CAP, and a method of screening for anticancer activity in a potential drug involving providing a cell that expresses a CA gene, contacting a tissue sample derived from a cancer cell with an anticancer drug candidate and monitoring the effect of the anticancer drug candidate on expression of the CA gene. The CAP proteins are useful for detecting cancer associated with expression of a CAP protein in a test cell sample and for screening for a bioactive agent capable of modulating the activity of a CAP protein. The CA nucleic acids are useful for diagnosing
   The invention relates to a method of determining susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain comprising identifying whether the individual has at least one polymorphism in a polymucleotide encoding at least one of the protein listed in the specification. The methods, composition and agent are useful for modulating the susceptibility of an individual to joint space narrowing and/or osteophyte development and/or joint pain that is associated with a disease, preferably osteoarthritis. The cell line and the non-human animal are useful for screening for an agent for diagnosing an individual having susceptibility to joint space narrowing and/or osteophyte development and/or joint pain. This sequence corresponds to the polynucleotide encoding a protein listed in the specification. (Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from WIPO at
  Human cancer-associated (CA)
  18-NOV-2004
   Sequence 190117 BP; 47446 A; 48907 C; 48857 G; 44888 T;
   Morris
   17-DEC-2002; 2002US-00322281
   15-DEC-2003; 2003WO-US040081
  15-JUL-2004.
  WO2004058146-A2
  Homo sapiens
  ds; cancer;
  Human; cancer-associated protein; CAP; cancer-associated gene; CA; gene;
  ABD33586;
  ABD33586 standard;
   ftp.wipo.int/pub/published_pct_sequences).
   (SAGR-)
   cancer
  99468
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACT 3115
   2004-499109/47
   human cancer associated protein encoded within open reading ncer associated gene, useful as targets for diagnosing cances
   45;
   B.
   SAGRES DISCOVERY
   Similarity
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACT 99512
   SEQ ID NO
   Malandro
  cytostatic.
   Conservative
  (first entry)
  DNA; 191584
   800; 182pp; English.
   X
O
   1.4%; Score 45;
100.0%; Pred. No.
   0;
  gene
   Mismatches
  HD07-118
   DB 10; 1
2.7e-10;
   0,
   Length 190117;
   Indels
   0
   U; 19 Other;
   cancer.
   0
   Gaps
   frame
   0
```

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muleic acid (I). Also described: (1) an expression vector comprising (I) (2) a host cell comprising (I) or the expression vector; (3) a comprising the defecting and nucleic acid; (4) an isolated cancer comprising that binds to the above polypeptide; (6) a hybridoma that composition composition that binds to the above polypeptide; (6) a hybridoma that composition that binds to the above polypeptide; (6) a hybridoma that composition the above antibody and a pharmaceutical composition comprising the above antibody and a pharmaceutical excipient; (8) a kit for detecting cancer cells, comprising the (monoclomal) antibody comprising the comprising the (monoclomal) antibody considered or absence of cancer cells in an individual; (10) a method for inhibiting growth of cancer cells in an individual; (11) a method for comprising the cancer cells in an individual; (12) an electronic library comprising the above polynucleotide or polypeptide, or their fragments; (13) methods of screening for anticancer activity or for
  Matches 45;
   Query Match
Best Local (
  cancer, involving determining the expression of a CA nucleic acid in a tissue. This sequence represents a human CA gene of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
  14-FEB-2003; 2003US-00367094.
14-MAR-2003; 2003US-003B8838.
23-SEP-2003; 2003US-00669920.
15-DEC-2003; 2003US-00737318.
   cancer; cancer associated nucleic acid; cancer associated gene;
cancer associated protein; CAP; cytostatic; vaccine; gene therapy;
lymphoma; leukaemia; human; gene; ds.
   The present invention describes an isolated cancer associated
  New isolated cancer-associated polynucleotides and polypeptides useful for diagnosing, preventing or treating cancers, especially lymphoma an leukemia, or in screening for agents that modulate cancer.
  Homo sapiens
  Human cancer associated gene genomic sequence SEQ ID NO:72
   ADR67026 standard; DNA; 191584 BP
   Sequence 191584 BP; 57287 A; 37750 C; 38021 G;
  Morris
   17-FEB-2004; 2004WO-US005000
  WO2004074321-A2
  18-NOV-2004
  (SAGR-)
  106736
  ftp.wipo.int/pub/published_pct_sequences
  3071 CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGACAAGACT
   2004-652915/63.
DB; ADR67028.
  16; SEQ ID NO 72; 166pp; English.
  DW, Malandro
  SAGRES
   Similarity
  CAAGATTGTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACT 106780
  Conservative
  (first
  DISCOVERY
  entry)
   1.4%;
  š
  0,
   Score 45;
Pred. No.
  Mismatches
   DB 13; I
2.7e-10;
  0,
   58526 T; 0 U; 0 Other;
   Length 191584;
  Indels
  3115
  lymphoma and
   0
   Ŝ
   Gaps
   0
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| a blostive spent capable of modulating the activity of a cot   100 kg    come made for description comes assessed with agreement of a cot   100 kg    come made for the composition of comes assessed with agreement of a cot   100 kg    come made for the composition and work of the comes as such in the composition of a cot   100 kg    come made for the composition and work of the composition of a cot   100 kg    come made for the composition and work of the composition of a cot   100 kg    come made for the composition and work of the composition of a cot   100 kg    come made for the composition and work of the composition of a cot   100 kg    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works are until the detection of the possess    composition and works     | E E E                              | 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 | F 1 1 1 1 1         |                                    | PPP       | F P P F        | 8 F F F F F                        | PPRX                                       | 8 X X 3                       | 2 2 2 2                                        | X X B                          | X Z Z X   | SXU E           | RES                      | 유 성             | <b>3</b> 00 1        | o so             | ៖<br>ខ្លួន ខ្លួន ខ | ន្តន្តន                                                       | នននន                                                                 |
|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------|---------------------------------------|---------------------|------------------------------------|-----------|----------------|------------------------------------|--------------------------------------------|-------------------------------|------------------------------------------------|--------------------------------|-----------|-----------------|--------------------------|-----------------|----------------------|------------------|--------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------|----------------------------------------------------------------------|
| justice of establiding the scriptly of a CVP, 131  proposed associating the scriptly of a CVP, 131  proposed associated substration of a polymorphism sensing of CVP, 131  proposed to the presence of an entitled in a term of the control of the proposed sensing of SVP, 131  proposed to the presence of an entitled in a term of the control of the proposed sensing of SVP, 131  proposed to the control of the contr | variation                          | variation                             | variation           | variation                          | variation | variation      | variation<br>variation             | Key<br>variation                           | guanine-nucleot Homo sapiens. | necology and<br>ne therapy; l<br>ngle nucleot: | zinc<br>tion;                  | 05        | 573             | JLT 189                  | 3071<br>06736   | i mi                 | Sequence 191584  | preventing and may also be use present sequence sequence, which                                                    | inhibiting the cytostatic acti                                | a bioactive age methods for det in a test cell serum sample; (       |
| Fit variation 7:109 k  | 2895<br>/*tag= j<br>/standard_name | ndard<br>g= i                         | J= g<br>ldard_<br>h | ndard<br>g= f<br>ndard             | e dard    | ှင့်ရှိ        | andard<br>ag= b<br>andard<br>5     | Location/Quali<br>207<br>/*tag= a          | ide exchange fa               | inte<br>poly                                   | and double PHD breast tumor; 6 |           | rd; DNA; 285300 |                          | TGTGCCACTGCACTO | 100.0%;<br>vative 0  |                  | treating cancer<br>d in screening<br>e represents a<br>is used in the                                              | expression of (vity, and can hand methods an                  | nt capable of mecting cancer as sample, or with 15) a method for     |
| Fit variation 7:109 k  | "Single                            | "Single                               | "Single             | "Single                            | "Single   | "Single        | "Single                            | fiers                                      | TE X TE                       | lasm; cytost<br>chromosome                     |                                |           | BP.             |                          | CAGCCTGGGCAACA  | Pred. No.<br>Mismatc | 7750 C; 38021 (  | exemplificati                                                                                                      | A gene in a ce<br>be used in vacc<br>e useful for d           | odulating the sesociated with the presence of treating can           |
| Fit variation 7:109 k  |                                    |                                       |                     |                                    |           |                | eotide polymorp<br>eotide polymorp |                                            |                               | . ஜ                                            | (DPF3)                         |           |                 |                          |                 | 0; Indels            | T; 0             | lymphoma and le<br>t modulate canc<br>ted gene genomi<br>on of the prese                                           | <ol> <li>The CA sequines, and in general sequines.</li> </ol> | activity of a C<br>expression of<br>of an antibody<br>cers; and (16) |
| retaction 1009 k reage 1 retaction 1009 k reage 1 retaction 1009 k reage 1 retaction 1000 meeting polymorphism retaction 1000 meeting poly |                                    |                                       |                     |                                    |           |                |                                    |                                            |                               | -                                              | mic DNA.                       |           |                 |                          | 780             | Gaps                 |                  | ent                                                                                                                | nences have<br>one therapy.                                   | AP; (14) a polypeptide in a test a method for                        |
| variation  /*tag= k /*tag= n / |                                    |                                       |                     |                                    |           |                |                                    |                                            |                               |                                                |                                |           |                 |                          |                 |                      |                  |                                                                                                                    |                                                               | <del></del>                                                          |
| tion  **trag= 1  /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 1 /*trag= 0 /*trag= ab /*trag= ab /*trag= ab /*trag= ac /*tra | H H H                              | 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 | 7777                |                                    | 1777      | 3777           | 7 7 7 7 7<br>1 1 1 1 1             | 7 7 7 7<br>1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 | FT                            | 9 7 7 7<br>1 1 1 1                             | F F F                          | F 17 17 1 | 7 7 7 X         | 1777                     | 1777            | 7 7 7 T              | 7777             | 17777                                                                                                              | F F F F                                                       | ** ** ** **<br>** ** ** **                                           |
| gs k  midard_name= "Single nucleotide polymorphism  property of the polymorphism  property of the property of the polymorphism  property of the property of the property of the polymorphism  property of the  | (* (                               |                                       |                     | 4 4                                | . 4       |                | 4 4                                |                                            | ď                             | rt .                                           |                                | רו ר      | 1 (             |                          | , 4             | rt                   |                  | ct ct                                                                                                              | et (                                                          |                                                                      |
| "Single nucleotide polymorphism                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | /*tag= ah<br>/standard_na<br>12604 | a aç                                  | ard<br>ard          | /*tag= ad<br>/standard_na<br>10823 | g= aq     | ndard<br>g= ab |                                    |                                            |                               | g= v<br>ndard<br>" "                           |                                |           |                 | /*tag= r<br>/standard_na |                 |                      |                  | /*tag= m<br>/standard_na<br>3642<br>/*tag= n                                                                       |                                                               | 3109<br>/*tag= k<br>/standard_na                                     |
| nucleotide polymorphism                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                    | , a                                   | a =                 | =                                  |           |                |                                    |                                            | =                             | 3                                              | =                              | II<br>=   | =               | n<br>=                   | }<br>=          | 11                   | a 11             |                                                                                                                    | 11                                                            |                                                                      |
| polymorphism                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | lingle                             | ingle                                 | ingle<br>ingle      | ingle                              | ingle     | single         | ingle                              | ingle<br>ingle                             | ingle                         | ingle                                          | Single                         | ingle     | ingle           | ingle                    | ingle           | ingle                | single<br>Single | dingle                                                                                                             | ingle                                                         | ingle                                                                |
| polymorphism                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    | nucleo                             | nucleo                                | nucleo              | nucleo                             | nucleo    | nucleo         | nucleo                             | nucleo                                     | nucleo                        | nucleo                                         | nucleo                         | nucleo    | nuclec          | nucleo                   | nucleo          | nuclec               | nucleo           | nucleo                                                                                                             | nucleo                                                        | nucleo                                                               |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                    |                                       |                     |                                    |           |                |                                    |                                            |                               |                                                |                                |           |                 |                          |                 |                      |                  |                                                                                                                    |                                                               |                                                                      |
|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                | olymorf                            | olymo                                 | olymor<br>olymor    | olymoı                             | olymo     | olymor         | olymor                             | olymor                                     | olymor                        | olymor                                         | olymor                         | olymor    | olymor          | olymor                   | olymor          | olymoz               | olymor           | olymo                                                                                                              | olymo                                                         | olymo                                                                |
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|                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                |                                    | 88                                    |                     |                                    |           |                |                                    |                                            |                               |                                                |                                |           |                 |                          | Sm .            | Sm.                  | 8 8              |                                                                                                                    | Sm.                                                           |                                                                      |

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   RESULT 190
  777777
CC in a subject. The method comparises detecting a mutation in the protein ctyrosine phosphatase 11 (PTPN11) gene in a subject, where the mutation cresults in increased PTPN11 gene is located on chromosome 12, more control. The human PTPN11 gene is located on chromosome 12, more control. The human PTPN11 gene is located on chromosome 12, more control. The human PTPN11 gene is located on chromosome 12, more control. The human PTPN11 gene is located on chromosome 12, more control. The human PTPN11 gene is located on chromosome 12, more control. The human PTPN11 gene is located to a kit for diagnosing Noonan syndrome, comprising an oligonucleotide that specifically hybridises to cor adjacent to a site of mutation of a PTPN11 protein, and control subject, comprising assessing the level of expression or activity of a PTPN11 protein in the test subject, and comparing it to the level of expression or activity of a pten in the test subject control is indicative of Noonan syndrome; (3) treating Noonan syndrome in a partial protein, and control is indicative of Noonan syndrome; (3) treating Noonan syndrome in control is indicative of Noonan syndrome; (3) treating Noonan syndrome in a solated to the control is indicative of Noonan syndrome; (3) treating Noonan syndrome; (4) an isolated prpN11 variant comprising a mutation resulting in increased level of PTPN11 activity; (5) an isolated cell comprising a control subject comprising a mucleic acid encoding the PTPN11 variant of (4), operatively associated with an expression control sequence; (6) an
   Query Match
Best Local S
Matches 45
   Gelb
  The present invention describes a method for diagnosing Noonan syndrome
   Claim 24; SEQ
  Diagnosing and treating Noonan a protein tyrosine phosphatase
  01-OCT-2001; 2001US-0326532P
   01-OCT-2002; 2002WO-US031290
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  Human PTPN11
  29-JAN-2004
   ADE86352
   variation
  variation
  WPI; 2003-381624/36
   10-APR-2003
   WO2003029422-A2
   Homo sapiens
   ADE86352
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   220046
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   "Single
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   (SNP) *
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| FT                    | 7 7 7<br>7 7 7                                    | 1777                                | 777                                 | 7 7 7<br>7 7 7                     | 7 7 7<br>7 7 7                       | F 1 T 1                            | F 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7                                                                                                                                 | PPT                                | P P P                                 | 8 8 X                                       | X                                                                    | <b>3 2 2</b>                                                                                                                                                             | 2222                                                                                                                                                                                                                                                                                                                                           | X E X                                                             | <b>33</b> 8          | ልጀ                   | RESU<br>ADO1                      | 유 &                                                     | % # O                                                                                                                                                                | ŝ                                                                       | នននា                                                                                                                                                                                                            | 8888                                                                                                                                                                                                                               |
|-----------------------|---------------------------------------------------|-------------------------------------|-------------------------------------|------------------------------------|--------------------------------------|------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------|---------------------------------------|---------------------------------------------|----------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------|----------------------|----------------------|-----------------------------------|---------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| exon 1                | /number= 4<br>/ intron 197501. 198676<br>/*tag= i | /number=<br>exon 197308.<br>/*tag=  | /number<br>intron 194626.<br>/*tag= | /number=<br>exon 194431.<br>/*tag= | /number=<br>intron 136954.<br>/*tag= | /number=<br>exon 136831.<br>/*tag= | /product="rivali" (198:246260246262,aa:Arg) /transl_except= (pos:246299246301,aa:Pro) /tansl_except= (pos:246299246301,aa:Pro) /tansl_except= (pos:246299246301,aa:Pro) | /number= 1<br>1235912<br>/*tag= b  | <pre>Key</pre>                        | Homo sapiens.<br>Synthetic.                 | kidney cancer; thyroid cancer; melanoma; leukaemia; human; gene; ds. | acute myeloid leukaemia; AML; juvenile myelomonocytic leukaemia; myelodysplastic syndrome; MDS; cancer; pre-cancerous condition; lung cancer: olorectal cancer: pancers. | protein tyrosine phosphatase gene 11; PTPN11; enzyme; protein tyrosine phosphatase gene 11 variant; PTPN11 variant; protein tyrosine phosphatase gene 11 variant; PTPN11 activity; cytostatic; haematologic disorder; mutation; increased PTPN11 activity; cytostatic; heuroprotective; PTPN11 modulator; acuite lumphoblastic lenkaemia; AII. | Human protein tyrosine phosphatase 11 gene sequence SEQ ID NO:33. |                      |                      | SULT 191<br>014076 standard, DNA. | 3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122 | Query Match 1.4%; Score 45; DB 10; Length 300000;<br>Best Local Similarity 100.0%; Pred. No. 2.6e-10;<br>Matches 45; Conservative 0; Mismatches 0; Indels 0; Gaps 0; | Sequence 300000 BP; 84671 A; 64420 C; 64260 G; 85849 T; 0 U; 800 Other; | aberrant expression and/or activity of the PTPN11 gene, specifically Noonan syndrome. The present sequence represents human PTPN11 genomic DNA, which is given in the exemplification of the present invention. | isolated nucleic acid encoding the PTPN11 variant of (4) isolated oligonucleotide which specifically hybridises tacid of (6). The methods and compositions of the present useful for diagnosing and treating a disorder associated |
| PD 21-MAY-2004.<br>XX | XX<br>PN WO2004041216-A2.<br>XX                   | exon 249938<br>/*tag= a<br>/number= | intron                              | exon 2                             | intron 2                             | exon 2                             | FT / rcag= z<br>PT / number= 13<br>PT intron 233289246256<br>PT / *tag= aa<br>PT / number= 13                                                                           | /*tag=<br>/number=<br>exon 233137. | /*tag= x<br>/number=<br>intron 232624 | /*tag= w<br>/number= 11<br>exon 23255623262 | /*tag= v /number=                                                    | /*tag= u<br>/number=<br>230588                                                                                                                                           | exon 226187 /*tag= t /number= /number= 22618                                                                                                                                                                                                                                                                                                   | FT 1NUION 222129225185 FT /*tag= 8 PT /number= 9                  | /*tag= r<br>/number= | FT exon 221970222128 | ٠ ۲                               | intron 2 / exon /                                       | /number=<br>exon 217057.<br>/*tag=<br>/number=                                                                                                                       | FT intron 200177. 217056 FT /*tag= m                                    | ٠ سر                                                                                                                                                                                                            | /*tag= j<br>/number=<br>intron 198794<br>/*tag= k                                                                                                                                                                                  |

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RESULT 192
ADJ12734
ID ADJ127
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AC ADJ127
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DT 20-MAY
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DT 20-MAY
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   control (II) has cytostatic and neuroprotective activity of PTPN11 protein and a carrier; (4) an isolated cc expression or activity of PTPN11 protein and a carrier; (4) an isolated cc cell (III) comprising a vector having (I), operatively associated with an CC expression control sequence; (5) an isolated nucleic acid encoding (I); cc and (6) characterising (M3) a cancer or pre-cancerous condition in a CC subject, which involves detecting a mutation in the PTPN11 gene in the cubic control. (I) has cytostatic and neuroprotective activities, and can be cused as a modulator of PTPN11 activity. (M2) is useful for treating a CC control. (I) has cytostatic and neuroprotective activities, and can be cused as a modulator of PTPN11 activity. (M2) is useful for treating a CC myeloid leukaemia (AM1), juvenile myelomonocytic leukaemia (AM1), acute myelodysplastic syndrome (MDS), in a patient. (M3) is useful for characterising a cancer or pre-cancerous condition in a subject, where creancer, kidney cancer, thyroid cancer, melanoma and leukaemia. The presents the human PTPN11 gene sequence, which is used control concer, melanoma and leukaemia is useful concer, in the exemplification of the present invention. The human PTPN11 gene is
  Matches
  Query Match
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   where the mutation corresponds to an amino acid substitution selected from Asn58Tyr, Gly60Val, Asp61Tyr, Asp61Val, Glu65Ly8, Phe71Ly8, Phe71Leu, Ala72Thr, Ala72Val, Ala72Asp, Glu76Ly8, Glu76Gly, Glu76Gly, Glu76Val, Ala72Val, Ala72Asp, Glu76Ly8, Glu76Gly, Glu76Val, Glu76Val, Glu76Val, Glu76Val, Ala72Val, Ala72Val, Ser502Pro, Gly503Arg, Gly503Arg, Gly503Arg, Thr507Ly8, Gln510Ly8, and combinations of them, in the human pyrN11 593 amino acid sequence of SRQ ID NO:2 (ADO14045). Also described: (1) characterising (M1) a haematologic disorder in a subject, which involves detecting a mutation in the PTPN11 gene in the subject, where the mutation results in an increased expression or activity of a PTPN11 protein encoded by the gene as compared to a control, or assessing the level of expression or activity of a PTPN11 protein in the test subject and comparing it to a control; (2) a kit (II) for diagnosing a haematologic disorder; (3) treating (M2) a haematologic disorder in a parient which involves adminieraring an accept that modulistics when
   The present invention describes an isolated protein tyrosine phosphatase \alpha == 11 (PTPN11) variant (I) associated with haematologic disorders, and
  Gelb BD,
   Claim 12; SEQ ID NO 33; 279pp; English.
  characterizing cancerous and precancerous conditions.
   05-NOV-2002; 2002US-0424170P.
  05-NOV-2003; 2003WO-US035349
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  Sequence 300001 BP;
   comprising a mutation resulting in an increased level of PTPN11 activity,
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  DNA fragment of a BAC clone that encodes a human secreted protein Seq588
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   105945 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 105989
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   protein tyrosine phosphatase gene 11 variant, useful for
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   UNIVERSITAETSKLINIKUM FREIBURG.
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This invention relates to novel polynucleotides encoding human secreted proteins. Specifically, it refers to the vectors, host cells, recombinant and synthetic methods for producing human polynucleotides, polypeptides and antibodies. Furthermore, it relates to screening methods to identify agonists and antagonists that can be used to inhibit or enhance the production and function of the secreted proteins. The present invention describes these compositions as useful for diagnosting, treating or preventing disorders such as cancer, haematopoietic diseases including anaemia and multiple myeloma, reproductive system disorders including prostatitis and inguinal hernia, musculoskeletal diseases including systemic lupus crythematosus and gout, cardiovascular disease including

arrhythmia and hypernatraemia,

mixed

Disclosure;

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NO 588; 286pp; English

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multiple myelowa; respondentive system disorder; prostatitis;
musculowa; reproductive system disorder; prostatitis;
musculowas; musculoskeletal disease; systemic lupus erythematosus;
KW gout; cardiovascular disease; arrhythmia; hypernatraemia; fetal disease;
KW detal alcohol syndrome; Down's syndrome; excretory disease;
KW urinary incontinence; renal disorder; neural; sensory disease;
KW verificational lung disease; endocrine disease; disbetes;
KW docupational lung disease; endocrine disease; disbetes;
KW glomerulonephritis; digestive disease; portal hypertension;
KW glomerulonephritis; digestive disease; portal hypertension;
KW wiritable bowel syndrome; epithelial disease; scleroderma;
KW miltipardic; anti-HIV; immunosuppressive; antianthritic;
KW antiasthmatic; anti-HIV; immunosuppressive; antianthritic;
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KW nephrotropic; uropathic; neuroprotective; antiparkinsonian; tranquilizer;
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   09-OCT-1997;
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09-OCT-1997;
New isolated nucleic acids and polypeptides, useful for diagnosing, treating, preventing or ameliorating diseases or disorders e.g. cancer, anemia, arthritis, asthma, inflammatory bowel disease or Alzheimer's
   30-OCT-2001;
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  09-OCT-1997;
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24-FEB-2000

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11-MAR-2000

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11-MAY-2000

07-JUN-2000

28-JUN-2000

07-JUL-2000

07-JUL-2000

11-JUL-2000

   antiarthritic, antiasthmatic, anti-HIV, immunosuppressive, antiinflammatory, antipsoriatic, antibacterial, osteopathic, dermatological, antipsoriatic, immunosuplator, antiarrhythmic, cardiant, nootropic, antilpemic, nephrotropic, uropathic, neuroprotective, antiparkinsonian, tranquilizer, antidiabetic, anabolic, hypertensive an vulnerary. This polynucleotide is a DNA fragment of a BAC clone that encodes a human secreted protein of the invention. NOTE: This sequence does not appear in the printed specification but has been obtained in electronic format from the US patent office at the following web site www.segdata.uspto.gov/sequence.html; Document ID: 20040010132.
  alcohol syndrome and Down's syndrome, excretory diseases including urinary incontinence and renal disorders, neural or sensory disease including Alzheimer's disease and meningitis, respiratory disease including emphysema and occupational lung disease, endocrine diseases including diabetes and glomerulonephritis, digestive diseases including portal hypertension and irritable bowel syndrome and connective tissue cepithelial diseases including scleroderma and epidermolysis bullosa. As such, there are various activites such as cytostatic, antianemic, antiarthritic, antiasthmatic, anti-HIV, immunosuppressive,
  Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis; ds.
  17-JAN-2001; 2001WO-US001354.
  09-AUG-2001.
   WO200157182-A2
  Homo sapiens
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   06-NOV-2001
  AAK67381;
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  2001-483426/52
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2000US-0251868P
  SEQ ID NO
   GENOME
  sc,
   SCI
   22193; 3071pp +
   Ruben
   SK.
```

Sequence

126

BP;

38

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3

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<u>و</u> 19 H

0 Ģ

0 Other;

```
example, they may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of (I) by expressing inactive proteins or to supplement the patients own production of (I). Additionally, (I) polynucleotides may be used to produce the secreted (I), by inserting the mucleic acids into a host cell and culturing the cell to express the protein. (I) proteins and polynucleotides may be used to prevent, diagnose and treat immune/haematopoietic-related diseases, especially cancers and cancer metastases of haematopoietic-derived cells. Ak64703 to AAK87694 represent human immune/haematopoietic antigen genomic sequences from the present invention. AAK874920 and AAK82169
  AAK54951 to AAK64702 encode the human immune/haematopoietic antigen (I) amino acid sequences given in AAM62170 to AAM91921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) proteins and polynucleotides may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate (I) expression. For
  human immune/hematopoietic antigen polypeptides, diagnosing and/or treating cancers and metastasis.
  Sequence Listing;
  English.
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28-JUN-2000;
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  07-NOV-2001
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   09-AUG-2001
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  2001WO-US001354.
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ID AAC218
XX
AC AAC218
XX
DT 06-OCT
XX
XX
DB Human
XX
KW Human;
KW gene t
  밁
  ঠ
   CC AAKS4951 to AAK64702 encode the human immune/haematopoietic antigen (I) CC amino acid sequences given in AAM82170 to AAM91921. (I) have cytostatic CC activity, and can be used in gene therapy and vaccine production. (I) CC proteins and polymucleotides may be used in the prevention, diagnosis and CC treatment of diseases associated with inappropriate (I) expression. For CC example, they may be used to treat disorders associated with decreased CC expression by rectifying mutations or deletions in a patient's genome CC that affect the activity of (I) by expressing inactive proteins or to CC supplement the patients own production of (I). Additionally, (I) CC polymucleotides may be used to produce the secreted (I), by inserting the CC protein. (I) proteins and polymucleotides may be used to prevent, CC diagnose and treat immune/haematopoietic-related diseases, especially CC cancers and cancer metastases of haematopoietic-derived cells. AAK64703 CC cancers and cancer metastases of haematopoietic antigen genomic CC sequences from the present invention. AAK34912 to AAK84950 and AAM82169 CC represent sequences used in the exemplification of the present invention.
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Best Local Similarity
Matches 44; Conserv
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Human; 5' \mathtt{EST}; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss.
   Disclosure;
   Nucleic acids encoding human immune/hematopoietic antigen polypeptides, useful for preventing, diagnosing and/or treating cancers and metastasis.
   Human secreted protein 5' EST, SEQ ID NO: 25910;
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2000US-0241B8BP.
2000US-0246AP.

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  Matches
   The present sequence is one of a large number of 5' ESTs derived from mRNAs encoding secreted proteins. No ORF has yet been conclusively identified within the present sequence. The 5' ESTs were prepared from total human RNAs or polyA+ RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3' untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5' ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5' UTR is rarely included. 5' ESTs are derived from mRNAs with intact 5' ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5' ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain upstream regulatory sequences and to design
   Human; digestive system antigen; gene therapy; cancer; ulcerative colitis; infection; Hirschsprung's disease; digestive system disorder; Meckel's diverticulum; ds.
  31-JAN-2000; 2000US-0179065P.
04-FEB-2000; 2000US-0180628P.
   New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for
  Dumas Milne Edwards J,
   17-JAN-2001; 2001WO-US001324.
   digestive system
   Human digestive system antigen genomic sequence SEQ ID NO: 4765.
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   expression and secretion vectors
  Claim 1; SEQ ID NO 25910; 71pp + Sequence Listing;
   diagnostic, forensic, gene therapy and chromosome mapping procedures
   26-FEB-1999;
  21-FEB-2000; 2000EP-00200610.
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  EP1033401-A2
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  02-AUG-2001.
   WO200155314-A2
  Homo sapiens
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   2000-500381/45
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  Similarity
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  (first
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  <u>.</u>
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Pred. No.
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20
Polynucleotides encoding digestive system antigens, useful for diagnosing, treating, preventing and/or prognozing disorders digestive system, particularly cancer and cancer metastases.
  2001-502630/55
   ξ
  HUMAN
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  2000US-0237039P.
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  GENOME SCI INC.
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rs of the
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Disclosure; SEQ ID

NO 4765; 986pp; English

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24-FEB-2000; 02-MAR-2000; 31-JAN-2000; 04-FEB-2000; 17-JAN-2001; 02-AUG-2001.

2000US-0179065P 2000US-0180628P 2000US-0184664P 2000US-0186350P

2001WO-US001351

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RESULT 197
AAS32121
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  Query Match
Best Local S
Matches 44
  cerebrovascular disorder; nervous system disorder; bacterial infection; fungal infection, viral infection; ocular disorder; endocrine disorder; gastrointestinal disorder; renal disorder; respiratory disorder; wound healing; skin aging; organ transplantation; tissue regeneration; anti-infertility.
  The present invention provides the protein and coding sequences of number of human digestive system antigens. These can be used in the diagnosis, treatment and prevention of bacterial or parasitic infections, appendicitis, Hirschsprung's disease, chronic colitis ulcerative colitis. The present sequence is a genomic DNA fragment
   Liver associated protein; human; mouse; rabbit; goat; horse; cat; dog; chicken; sheep; immunosuppreqsive; antiarthritic; vasotropic; antiproliferative; cytostatic; cardiant; neuroprotective; cerebroprotective; nootropic; antibacterial; virucide; fungicide; cancer ophthalmological; vulnerary; gene therapy; autoimmune disease; neoplasm; hyperproliferative disorder; breast; liver; cardiovascular disorder; ds;
   WO200155355-A1
   Human liver associated genomic DNA #295
   04-DEC-2001
  AAS32121;
  AAS32121
   Sequence 182 BP; 48
  encoding a digestive system antigen of the invention
   3078 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGAGCTCTGTCT 3121
   sapiens.
   139 GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT 182
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   Similarity
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   Conservative
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   A; 44 C; 55 G; 35 T; 0 U; 0 Other;
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Pred. No.
   Mismatches
   DB 4,
a. 1e-09;
0;
   Length
   Indels
   182;
   colitis or
  ç,
  the
   0
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Sequences AAS31827-AAS32182 represent genomic DNA molecules, which encode the liver associated polypeptides of the invention. Liver associated CC the liver associated polypeptides and their associated polypuncleotides are useful in the CC diagnosis, treatment and prevention of various types of disorders in e.g. CC humans, mice, rabbits, goats, horses, cats, dogs, chickens or sheep. A CC pathological condition can be determined by detecting the presence or CC absence of a mutation in a liver associated polynucleotide. The treatable disorders include autoimmune diseases such as rheumatoid arthritis, CC cardiovascular disorders such as cardiac arrest, cerebrovascular cardiovascular disorders such as cardiac arrest, cerebrovascular CC disorders such as cardiac arrest, cerebrovascular associated by bacteria, viruses and fungi, coular disorders such as corneal infection, endocrine disorders such as corneal infection, endocrine disorders such as CC premature labour and infertility, gastrointestinal disorders such as CC cohn's disease, renal disorders such as glomerulonephritis and CC respiratory disorders such as asthma and pleurisy. The polypeptides can CC calso be used to aid wound healing, to prevent skin aging due to sumburn, to maintain organs before transplantation, to regenerate tissues and in chemotaxis. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly
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08-NOV-2000
17-NOV-2000
17-NOV
   Isolated nucleic acid used in preventing, to particularly cancer of
  Rosen
   Claim
   (HUMA-)
  ς,
   1,
  HUMAN
   SEQ
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  2000US-0246526P

2000US-0246528P

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2000US-0251939P

2000US-0251868P

2000US-0251868P

2000US-0251868P

2000US-0251869P

2000US-0251869P

2000US-025199PP

  IJ
  GENOME
   ð
  sc,
   cid molecule e
, treating or
r of the liver
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  526pp;
   X.
  English.
   encoding a human liver related ameliorating disorders of the
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liver

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01-SEP-2000

   Human; liver antigen; liver disorder; hepatic disorder; infection; hepatitis; viral; parasitic; bacterial; fungal; inflammatory condition; cirrhosis; granulomatous hepatitis; toxin damage; drug damage; autoimmune disease; wilson's disease; primary biliary cirrhosis; neoplastic disorder; cancer; tumour; portal hypertension; gastrointestinal disorder; hepatitis; drug screening; gene therapy; chromosome mapping; forensic analysis; antibody preparation; hepatotropic; cyrostatic; antiinflammatory; virucide; antibacterial; fungicide; parasiticide; antidote; immunosuppressive; gene; ds.
   07-JUL-2000;
  31-JAN-2000;
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   11-APR-2002
  US2002042096-A1
   Human liver antigen HHLAB49 genomic sequence, SEQ ID NO:597
  24-JUL-2002
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2000US-022576P.
2000US-0225785P.
2000US-0235834P.
2000US-0234274P.
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  2001US-00764887.
  (first entry)
  2000US-0180628P
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  182
  Score 44;
; Pred. No.
  0; Mismatches
  DB 5;
1e-09;
  0;
  Length 182;
  Indels
  0
  Gaps
  0
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δ

Query Match Best Local S Matches

Local Similarity

1.4%;

Score 44; Pred. No. 55 G; 35 T;

B 1e-09;

Length 182 Indels

0,

Gaps

0

44;

Conservative

<u>.</u>

Mismatches

Sequence

182

BP;

48 ₽,

44 C;

0 U; 0 Other; 6; 0

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```
CC liver antigen polymucleotides and polypeptides in diagnosing, treating, conclude viral infections (e.g., cytomegalovirus, Epstein-Barr virus, inpatitis B virus and hepatitis C virus), parasitic confections (e.g., Clonorchis sinensis, Echinococcus granulosus and confections (e.g., clinorchis sinensis, Echinococcus granulosus and confections (e.g., clinorders that may be treated include inflammatory conditions (e.g., clinorders that may be treated include inflammatory conditions (e.g., concolistic disorders (e.g., wilson's disease, primary biliary cirrhosis), concopiastic disorders (e.g., adenomas, haemangiomas and hepatocellular concopied culcers, gastritis and peritoneal disease). Liver antigen copypeptides and polymucleotides may also be used in screening for compounds which modulate liver antigen expression or activity. The colypeptides may be used as molecular weight markers or to prepare colypeptides may be used as molecular weight markers or to prepare colypeptides may be used as molecular weight markers or to prepare colypeptides may be used as molecular weight markers or to prepare contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful in disease diagnosis, drug targeting and phenotyping. Contibodies useful diagnosis, drug targeting did not form part of the printe
  29-SBP-2000
29-SBP-2000
29-SBP-2000
29-SBP-2000
02-OCT-2000
  The invention relates to 145 novel human liver antigens (ABP40831-ABP40975) and to cDNAs encoding them (ABN90036-ABN90180), and also encompasses polypeptides 90% identical and polymucleotides 95% identical to the sequences of the invention. The invention additionally relates to recombinant vectors and host cells comprising human liver antigen polynucleotides, antibodies against human liver antigens, and the use of
  New nucleic acid encoding human liver antigens, useful for diagnosis, treatment and prevention of e.g. hepatitis and hepatic cancer, also related polypeptides and antibodies.
   Disclosure; SEQ ID NO 597; 181pp; English.
   Rosen
  WPI; 2002-381944/41.
  (ROSE/)
   08-DEC-2000;
  17-NOV-2000;
  (BARA/)
   CA, Ruben SM,
  ROSEN
RUBEN
  BARASH S
  2000US-023636PP.
2000US-0236368PP.
2000US-0236369PP.
2000US-0236370PP.
2000US-0237037PP.
2000US-0237037PP.
2000US-0237039PP.
2000US-0237040PP.
2000US-0237040PP.
2000US-0240960PP.
2000US-0241785PP.
2000US-0241809PP.
2000US-0251868PP.
2000US-0251868PP.
  S C M
   Barash SC;
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| 14-AUG-2000<br>18-AUG-2000<br>22-AUG-2000<br>22-AUG-2000<br>23-AUG-2000<br>30-AUG-2000<br>01-SEP-2000<br>01-SEP-2000<br>01-SEP-2000<br>05-SEP-2000<br>05-SEP-2000 | PR 07-JUL-2000; 2000US-0216880P. PR 11-JUL-2000; 2000US-0217487P. PR 11-JUL-2000; 2000US-0217496P. PR 14-JUL-2000; 2000US-0218290P. PR 26-JUL-2000; 2000US-0228963P. PR 26-JUL-2000; 2000US-0220964P. PR 14-AUG-2000; 2000US-0224518P. PR 14-AUG-2000; 2000US-022513P. PR 14-AUG-2000; 2000US-0225214P. PR 14-AUG-2000; 2000US-0225214P. PR 14-AUG-2000; 2000US-0225266P. PR 14-AUG-2000; 2000US-0225266P. PR 14-AUG-2000; 2000US-022526P. PR 14-AUG-2000; 2000US-022526P. PR 14-AUG-2000; 2000US-0225270P. PR 14-AUG-2000; 2000US-0225270P. PR 14-AUG-2000; 2000US-02252757P. PR 14-AUG-2000; 2000US-02252757P. | 14-FEB-2002<br>31-JAN-2000<br>04-FEB-2000<br>24-FEB-2000<br>16-MAR-2000<br>17-MAR-2000<br>17-MAR-2000<br>19-MAY-2000<br>07-JUN-2000<br>30-JUN-2000<br>30-JUN-2000<br>07-JUL-2000 | cytostatic; gynaecological; viral; fungal; bacterial; paraeitic infection; cirrhosis; Wilson's disease; gastrointestinal disorder; pancreatic; gallbladder; immune; blc hyperproliferative; cardiovascular; respiratory; musculoskelete neurological; endocrine; reproductive system; developmental; ir human; ds.  Homo sapiens.  US2003077602-A1. | 150                                    |
|-------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------|
| ***************************************                                                                                                                           | ***************************************                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          | ######################################                                                                                                                                           | 2                                                                                                                                                                                                                                                                                                                                                   | ###################################### |
| 000000000000000000000000000000000000000                                                                                                                           | 20 - OCT - 2000 20 - OCT - 2000 01 - NOV - 2000 08 - NOV - 2000                                                                                                                                                                                                                                                  |                                                                                                                                                                                  |                                                                                                                                                                                                                                                                                                                                                     |                                        |
| 200000000000000000000000000000000000000                                                                                                                           | 2000US-0241809P. 2000US-02446174P. 2000US-0246474P. 2000US-0246475P. 2000US-0246476P. 2000US-0246478P. 2000US-0246478P. 2000US-0246523P. 2000US-0246525P. 2000US-0246528P.                                                                                                                                                                                                                                            |                                                                                                                                                                                  |                                                                                                                                                                                                                                                                                                                                                     |                                        |

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RESULT 200
AAK87671/c
ID AAK876
XX
AC AAK876
XX
AC AAK876
XX
OT-NOV
  문
  Ś
  Query Match
Best Local Similarity
Matches 44; Conserv
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17-NOV-2000;
17-NOV-2000;
17-NOV-2000;
01-DEC-2000;
05-DEC-2000;
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06-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
08-DEC-2000;
  osteopathic, mortopic, antiparkinsonian, anticonvulsant, neuroleptic, vasotropic, cytostatic and gynaecological activities. The polypeptides and polynucleotides of the invention may be useful for diagnosis, detection, treatment and/or prevention of disorders of the liver such as viral, fungal, bacterial or parasitic infections, cirrhosis, Wilson's disease, gastrointestinal disorders, pancreatic disorders, gallbladder disorders, immune disorders, blood related disorders, hyperproliferative disorders, cardiovascular disorders, respiratory disorders, musculoskeletal system disorders, neurological diseases, endocrine disorders, reproductive system disorders or developmental and inherited disorders. The current sequence is that of the human liver-related genomic DNA of the invention. The current sequence is not shown within the specification per se but was obtained electronically from the USPTO
 07-NOV-2001
                                    AAK87671;
   AAK87671 standard;
   the specification per se web-site.
  The invention relates to a novel isolated, liver related polypeptide. The polypeptide of the invention demonstrates virucide, fungicide, antiparasitic, hepatotropic antiinflammatory, cytostatic, litholytic, antirheumatic, antiarthritic, neuroprotective, antidiabetic, anticoagulant, thrombolytic, antiarteriosclerotic, cardiant, haemostatic, antiarrhythmic, ophthalmological, antiarteriosclerotic, vasotropic, antiarrhythmic, ophthalmological, antiarteriosclerotic, vasotropic,
   Disclosure; SEQ ID NO
   New liver related polypeptide, useful for diagnosis, treatment and/or prevention of liver, gastrointestinal, pancreatic, immune, blood related, endocrine, reproductive, hyperproliferative or reproductive disorders.
   WPI; 2003-765398/72.
   (HUMA-) HUMAN GENOME
   7-NOV-2000;
7-NOV-2000;
7-NOV-2000;
   3078
  139
  ξ
  GTGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCT 3121
  GTGCCACTGCAGCCTGGGCAACAGAGCAAGACTCTGTCT 182
  Ruben
  2000US-0249218P.
2000US-024924SP.
2000US-024926SP.
2000US-024929SP.
2000US-024929SP.
2000US-024929SP.
2000US-024929SP.
2000US-0251930SP.
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 (first entry)
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   SCI INC
  597; 181pp; English.
  Barash
   307
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Human; immune; haematopoietic; immune/haematopoietic antigen; cancer; cytostatic; gene therapy; vaccine; metastasis; ds.
  Human
  immune/haematopoietic antigen genomic sequence
  SEQ ID NO: 42483
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Homo sapiens.

WO200157182-A2

31-JAN-2000; 04-FEB-2000; 24-FEB-2000; 02-FEB-2000; 02-MAR-2000; 16-MAR-2000; 17-MAR-2000; 18-APR-2000; 17-JAN-2001; 2000US-0179065P. 2000US-0180628P. 2001WO-US001354

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2000US

19-MAY-2000
17-JUN-2000
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11-JUL-2000
11-JUL

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2000US-0246611P
2000US-024652P
2000US-024921P
2000US-024921P
2000US-024921P
2000US-024921P
2000US-025939P
```

```
cc amino acid sequences given in AAM82170 to AAM81921. (I) have cytostatic activity, and can be used in gene therapy and vaccine production. (I) cc proteins and polynucleotides may be used in the prevention, diagnosis and ct treatment of diseases associated with inappropriate (I) expression. For cexample, they may be used to treat disorders associated with decreased cexpression by rectifying mutations or deletions in a patient's genome ct that affect the activity of (I) by expressing inactive proteins or to complement the patients own production of (I). Additionally, (I) complement the patients own production of (I). Additionally, (I) concleic acids into a host cell and culturing the cell to express the complement call in the proteins and polynucleotides may be used to prevent, (I) contein. (I) proteins and polynucleotides may be used to prevent, (I) concers and cancer metastases of haematopoietic related diseases, especially concers and cancer metastases of haematopoietic derived cells. AAK64703 to AAK687694 represent human immune/haematopoietic antigen genomic compresent sequences used in the exemplification of the present invention. AAK54942 to AAK54950 and AAM82169 represent sequences used in the exemplification of the present invention.
 뭐
                             ঠ
   Query Match
Best Local Similarity
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08-DEC-2000; 2000US-0251999P.
11-DEC-2000; 2000US-0254997P.
05-JAN-2001; 2001US-0259678P.
   Sequence 307 BP; 53
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  Rosen
   AAK54951 to AAK64702
   Disclosure;
  (HUMA-) HUMAN GENOME
  δ
74
  acids encoding
                         TGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC 3122
 TGCCACTGCACTCCAGCCTGGGCAACAGAGCAAGACTCTGTCTC
   preventing,
  Barash SC,
   1.4%; So ilarity 100.0%; I Conservative 0;
   SEQ ID NO 42483; 3071pp + Sequence Listing; English
   A; 93
  SCI
   encode the human immune/haematopoietic antigen (I) encode the human immune/haematopoietic antigen (I)
   human immune/hematopoietic diagnosing and/or treating
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   C; 69 G;
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   92
   Η,
   DB 4; Lo
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   ď
   0 Other;
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   antigen polypeptides, cancers and metastasis.
   31
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Gaps